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FILE 'BIOSIS' ENTERED AT 19:01:24 ON 21 JAN 2003 COPYRIGHT (C) 2003 BIOLOGICAL ABSTRACTS INC.(R)

=> s p53 and antisense L1 4478 P53 AND ANTISENSE

=> s l1 and splice acceptor site 78 L1 AND SPLICE ACCEPTOR SITE L2

=> dup rem 12 PROCESSING COMPLETED FOR L2 78 DUP REM L2 (0 DUPLICATES REMOVED)

=> s 13 and morpholino 2 L3 AND MORPHOLINO L4

=> d 14 ibib abs tot

ANSWER 1 OF 2 CAPLUS COPYRIGHT 2003 ACS 2001:816897 CAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER:

135:353717

TITLE:

Splice-region ***antisense*** oligonucleotide composition and targeting the mRNA splicing

INVENTOR(S):

Iversen, Patrick L.; Hudziak, Robert Avi Biopharma, Inc., USA

PATENT ASSIGNEE(S): SOURCE:

PCT Int. Appl., 53 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO. KIND DATE APPLICATION NO. DATE wo 2001-US14410 20010504 wo 2001083740 Α2 20011108

W: AU, CA, JP, KR RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL,

PT, SE, TR PRIORITY APPLN. INFO.:

US 2000-202376P P 20000504

Antisense compns. targeted against an mRNA sequence for a selected protein, at a region having its 5' end from 1 to about 25 base pairs downstream of a normal splice acceptor junction in the preprocessed mRNA, are disclosed. The ***antisense*** compd. is Rnase-inactive, and is preferably a phosphorodiamidate-linked ***morpholino*** oligonucleotide. Such targeting is effective to inhibit natural mRNA splice processing, produce splice variant mRNAs, and inhibit normal expression of the protein.

ANSWER 2 OF 2 USPATFULL

ACCESSION NUMBER:

2000:57750 USPATFULL

TITLE:

Chimeric oligonucleoside compounds

INVENTOR(S):

Arnold, Jr., Lyle J., Poway, CA, United States

PATENT ASSIGNEE(S):

Giachetti, Cristina, Solano Beach, CA, United States Lebedev, Alexandre V., San Diego, CA, United States Genta Incorporated, Lexington, MA, United States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION:

us 6060456 20000509

APPLICATION INFO.: RELATED APPLN. INFO.: US 1997-960111 19971027 (8)
Continuation of Ser. No. US 1995-481637, filed on 7 Jun 1995, now abandoned which is a continuation of Ser. No.

US 1994-238177, filed on 4 May 1994, now abandoned which is a continuation of Ser. No. US 1994-233778, filed on 26 Apr 1994, now abandoned which is a continuation of Ser. No. US 1993-154013, filed on 16

Nov 1993, now abandoned which is a continuation of Ser. No. US 1993-154014, filed on 16 Nov 1993, now abandoned Utility

DOCUMENT TYPE: FILE SEGMENT:

Granted Riley, Jezia

PRIMARY EXAMINER: LEGAL REPRESENTATIVE:

Knobbe, Martens, Olson & Bear LLP.

NUMBER OF CLAIMS:

38

EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

15 Drawing Figure(s); 12 Drawing Page(s)

LINE COUNT:

5081

CAS INDEXING IS AVAILABLE FOR THIS PATENT. AB

Chimeric oligonucleoside compounds, and methods of preparing and formulating the same, are disclosed. The compounds and compositions are useful in activating RNaseH-mediated cleavage of target ribonucleic acid sequences, and in treating disease conditions relating to such sequences.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

=> d 14 2 kwic

SUMM

L4 ANSWER 2 OF 2 USPATFULL

The present invention relates to ***antisense*** SUMM oligonucleoside compounds containing modified internucleoside linkages, and optionally other structural modifications. The compounds are capable of hybridizing

to target nucleic.

Considerable attention has been directed in recent years to the design SUMM ***antisense*** nucleic acid oligomers for use in studying, treating and diagnosing conditions attributable to endogenous or foreign nucleic acid sequences in living organisms. For example, it is now well known that a nucleic acid oligomer having suitable ***antisense*** complementarity to a target mRNA can hybridize to the target mRNA and, in some cases, disrupt translation of the mRNA. The ***antisense*** approach presents great promise for the eventual therapeutic treatment of disease conditions caused by foreign (e.g., viral) genetic material,

However, despite the great promise of the ***antisense*** approach a number of challenges still remain. First, ***antisense*** compounds are generally subject to degradation in the cellular milieu SUMM approach, due to endogenous endo- and exonucleases. While a number of modified ***antisense*** structures have been described having improved

resistance to nuclease degradation, further improvements are desirable in order to increase the potency and half-life of the compounds. Second, it is generally required that an ***antisense*** compound have a high specificity toward the intended target nucleic acid so as to avoid disruption of activity of unintended native sequences. Although a number of researchers have described approaches designed to increase the binding affinity of an ***antisense*** compound to a target sequence, very few results have been reported with respect to structural

refinements which avoid disruption of.

mRNAs involves forming a duplex hybrid between the target mRNA and an ***antisense*** strand, followed by cleavage of the target mRNA by an endogenous RNaseH. See Dash, P., et al, Proc. Natl. Acad...
Nucleases (Linn & Roberts, eds.), Cold Spring Harbor Laboratory (1982), at 212. As a result, one putative requirement of the ***antisense***
RNaseH cleavage approach is that at least some of the nucleosides of the ***antisense*** nucleic acid strand must have characteristics in

common with deoxyribonucleotides (as opposed to ribonucleotides) particularly, the absence of a polar group on the 2'-position of the

```
additional requirement that at least some of the sugar groups in the ***antisense*** compound must be in a 2'-endo (.beta.) conformation as found in deoxyribonucleosides, as opposed to the 3'-endo (.alpha.)
                 conformation found.
                 It has further been reported that various 2'-position substituents (e.g., 2'-O-alkyl and 2'-fluoro) will render the substituted portion of an ***antisense*** strand non-activating to RNaseH, even though
 SUMM
                 binding affinity toward the target nucleic acid is increased. Inoue, H.,
                et al., FEBS. . . achieve efficient activation of mammalian (HeLa) RNAseH, and that this 2'-deoxy segment (if accompanied by 2'-substituted residues in the same ***antisense*** compound) must be centered in
                 the oligomer sequence in order to achieve efficient RNaseH activation in
                 vitro or expression inhibition.
                 Another reported requirement of the ***antisense***
 SUMM
                                                                                                                                             RNaseH cleavage
                approach is that, in order to achieve RNaseH activation, at least one portion of the internucleoside "backbone" of the ***antisense***
                compound must include charged (anionic) phosphorus-containing linkage groups. Cook, P. D., PCT Publication No. WO 93/13121 (1993), at 18. In studies of chimeric ***antisense*** compounds including both
                methylphosphonate (uncharged) and phosphodiester or phosphorothioate (charged) linkages, Agrawal, et al. reported that the minimum number of.
                . . . exonucleases. A variety of alternative linkage groups, some of which are nuclease-resistant, have been developed or proposed for use with ***antisense*** compounds. Among these are charged linkage
SUMM
                groups such as phosphorothioate, phosphorodithioate, phosphoroselenate and phosphorodiselenate linkers. In general, deoxyribonucleoside ***antisense*** oligomers containing these non-natural linkage groups tend to have lower binding affinity toward complementary RNA target strands than the corresponding phosphodiester-linked ***antisense***
                strands than the corresponding phosphodiester-linked ***antisense***
oligomers, although higher affinity may be achieved where the
    ***antisense*** strand comprises ribonucleosides or 2'-substituted
                 ribonucleosides (rather than deoxyribonucleosides). See Metelev, v. &
               Agrawal, S., PCT Publication No. WO 94/02498. . . Padmapriya, A. & Agrawal S., PCT Publication No. WO 94/02499 (1994). Non-phosphorus-based linkage groups have also been reported, including peptide, ***morpholino***, ethylene glycol, amide, and other linkers. See Reynolds, M. A., et al., PCT Publication No. WO 92/02532 (1992); Cook, P. . . lower binding affinity (compared to phosphodiester linkages) toward complementary RNA target strands, at least in the case of linked 2'-unsubstituted ***antisense*** nucleotides, and particularly in the presence of salt ions
                the presence of salt ions.
SUMM
                Various workers have attempted to identify combinations of linkage
                groups and/or structural modifications for
                                                                                                                    ***antisense***
                                                                                                                                                             oligomers
               that might lead to improved RNaseH activation, binding affinity, nuclease resistance and/or target specificity. Thus, Cohen, et al. have reported improved half-life for ***antisense*** and non-
***antisense*** oligodeoxyribonucleotides containing at least one
               ***antisense*** oligodeoxyribonucleotides containing at least one phosphorothioate linkage located, for example, at either terminus of the compound, or throughout the compound. . . . RNaseH cleavage activation
                compound, or throughout the compound. . . RNaseH cleavage activation reportedly required retention of at least four, and preferably at least seven, contiguous phosphodiester linkages in the ***antisense***
               oligomer. The preferred compounds contained at least 10, and preferably at least 15, nucleotides, the majority of which were
                phosphodiester-linked..
               Giles & Tidd have reported that the target specificity of an

***antisense*** oligomer can be improved by the use of a chimeric
structure comprising terminal methylphosphonodiester sections separated
SUMM
                by a central RNaseH-activating.
SUMM
                                  was reportedly localized to a site (or sites) on the target
               corresponding to the non-substituted (i.e., deoxyribonucleotide) portion of the ***antisense*** compound. Single-site cleavage was reportedly optimized by use of a tetradeoxyribonucleotide segment located centrally
               in the compound between two 2'-substituted.
               The present invention relates to improved RNaseH-activating
***antisense*** oligonucleoside compounds containing selectively
SUMM
               modified internucleoside linkages, and optionally other structural modifications. The compounds exhibit improved target specificity and
               potency compared to other RNaseH-activating ***antisense*** compounds. They are useful both in vivo and in vitro in reducing or eliminating the translation of target mRNA sequences,. . . .
SUMM
               In another aspect, the present invention provides chimeric structures
               for ***antisense*** oligonucleoside compounds that maximize activity while maintaining the ability to effect selective RNaseH-mediated cleavage of the intended target strand. These.
SUMM
               In another aspect, the present invention includes imp roved
```

diagnosing diseases or other conditions in living organisms attributable to the expression of endogenous. Consider first the challenge of achieving high target specificity with an ***antisense*** cleavage compound. Mammalian cells typically contain an RNA population comprising about 3.times.10.sup.7 ribonucleotides. By assuming a statistically random distribution of. DETD . . . mismatches increases. If the K.sub.A for a given mismatch duplex is sufficiently high as to allow appreciable hybridization of an ***antisense*** oligomer to a mismatched target, then unintended and DETD undesirable cleavage of the mismatched target can result. Take, for example, the case of a one-base mismatch between a 12-to-18 nucleoside ***antisense*** oligomer and an unintended mismatch RNA sequence. The present inventors have ascertained that the K.sub.A for the correct "match" hybridization. **DETD** and polyadenylation sites), inhibition of protein production DETD can be achieved prior to the translation process by suitable hybridization of an ***antisense*** oligonucleoside, and ribosomal displacement of the hybridized oligomer generally does not occur. As a result, oligonucleosides having higher binding affinities. . . . reviewed by Jaeger et al., Annual Reviews in Biochemistry 62, 255-287 (1993). Another approach is to utilize two or more ***antisense*** compounds in tandem, at least one of which is a chimeric oligonucleoside of the invention, which ***antisense*** DETD compounds have nucleoside base sequences selected to hybridize to adjacent regions in a secondary-structured mRNA target region. It is known that adjacently-hybridizing ***antisense*** compounds may be used to disrupt secondary structure of RNA molecules and thus to enhance the effective K.sub.A 's of. As discussed above in the background section of this disclosure, a number of workers in the ***antisense*** field have reported various and disparate efforts to increase binding affinity of ***antisense*** DETD oligonucleosides, to optimize RNaseH activation, to improve nuclease resistance, and to improve target specificity. It will be seen in light. As is also explained above, one putative requirement of mammalian RNaseH activation is that the ***antisense*** compound must have a sequence DETD of at least four or five consecutive charged (anionic) internucleoside DETD gene expression comprising one or more segments with methylphosphonate internucleosidyl linkages enhanced for the R.sub.p configuration which DETD originally derived from the coupled dimeric units). The remaining internucleosidyl linkages comprise non-phosphorate internucleosidyl linkages, such as phosphodiester, phosphorothioate, phosphorodithioate, ***morpholino***, phosphoramidite, phosphorofluoridate, boranophosphate, formacetal, silyl or other non-phosphonate internucleosidyl linkages. DETD that the sequence of nucleoside bases in the present chimeric RNaseH-activating compounds, and also the sequence of bases in RNaseH-activating ***antisense*** compounds generally, can RNaseH-activating ***antisense*** compounds generally, can be selected in a manner described herein to provide RNaseH-activated cleavage that is highly specific to disease. . . caused by sin (particularly single-base) mutations, allele differences or other caused by singular anomalies in genetic sequence. In particular, the inventors have discovered ***antisense*** oligonucleoside constructions that are capable of discriminating a single base difference in a target RNA In this regard, ***antisense*** oligonucleosides have reportedly been used to target a wide variety of mRNAs associated with cancer and other diseases. See, for. . . Science, 261, 1004-1012 (1993); Uhlmann, E. and Peyman, A., Chemical Reviews, 90, 544-584 (1990). One challenge in the application of ***antisense*** oligonucleosides as therapeutic agents is the need to discriminate a single base difference between target and non-target sequences. A number. . . for example, have been identified which differ from their normal counterparts by only a single base change, including the RAS, ***p53***, src and erbB-2 DETD a single base change, including the RAS, ***p53***, src and erbB-2 oncogenes. Single base changes are also associated with some inherited genetic diseases such as Lesch-Nyhan syndrome (Fujimura,. . . Ideally, and within the constraints described in the specification above, ***antisense*** oligonucleosides should have high affinities above, ***antisense*** oligonucleosides should have high affinities with their target sequences, so that small amounts of oligonucleoside can be used for treatment.... DETD DETD . . oligomers can be achieved by carefully defining the position of

```
***antisense*** oligomer. Particularly, the oligomer is selected, first, to have a base sequence that is complementary to a target region
           . . . hypotonic dounce lysis in 5.times. the packed cell volume. It was buffered to pH 6.0 by adding 0.4 mL of 2-(N- ***morpholino*** ) ethanesulfonate (MES, 0.5 M solution, pH 6.0) to 3.6 mL of cell lysate
DETD
           on ice and mixing with mild agitation..
           Inhibition of Protein Synthesis in a Cell Culture With Chimeric

***Antisense**** Oligomers Targeted to a Non-Eukaryotic Reporter Gene,
DETD
           Chloramphenicol Transferase
DETD
           The following example shows the ability of chimeric
                                                                                              ***antisense***
           oligomers to selectively inhibit protein synthesis in a eukaryotic cell
           culture system. COS-7 cells were transiently transfected with plasmids encoding either a target reporter gene or a control non-target reporter gene. These cells were then treated with various chimeric

***antisense*** or control oligomers and then assayed for the
           expression of the reporter genes. . . . tct gca 3'
DETD
   XV-6 SEQ ID NO:9,
    24mer, all phosphorothioate:
       cac tca atc aat gac tag tct gca 3'
***Splice*** ***acceptor***
                                                                 ***site***
                                                                                     oligomers:
   3265-1 SEQ ID NO:19,
24mer, (MP(R.sub.p)/DE)(PS/DE)(MP(R.sub.p)/DE):
    5' ccc tga ga(g aga g)ag aga ggt tcg 3'
    3266-1 SEQ ID.
DETD
           Anti-Splice Site Oligomers Versus pG1035 and pG1036 (splicing inhibition
                   ***antisense***
                                             oligomers):
                         splice site acceptor oligomer 3387-1 and lengthening the PS
DETD
           center from five to seven continuous phosphorothioate backbone linkages increases the ***antisense*** activity against the ***splice***

***acceptor*** ***site*** target significantly but does not
           increase non-specific activity against the control target.
                    . all-PS oligomers and control chimeras. Both target-specific and
DETD
           oligomer-specific controls were included, demonstrating that the results are based on sequence-specific ***antisense*** effects.
           . . . of an oligonucleoside and/or in the target mRNA. The chimeric compounds listed below (see also Example 41) were assayed for ***antisense*** activity against both the pG1040 (UCAT) target and the pG1042 (UCAT) 4-base mismatch control. The oligomer sequences were as
DETD
           follows.
DETD
           pG1040 (UCAT) target mRNA and
                                                            ***antisense***
                                                                                       oligomers:
              +1 +4
                                                                 +27
                 .vertline.
                                   .vertline.
                                                                                                 .vertline.
                 Met Glu Lys Lys Ile Ser Gly Tyr Thr ... ...
   DETD
                                        activity was assayed against both pG1041 (UCAT) and
           pG1042 (UCAT) using procedures as generally described in Example 41,
                                         It was demonstrated that mismatches in the
           phosphorothioate core and the position of the core in chimeric oligomers greatly affected ***antisense*** activity. The following table sets
           forth the percentage of gene expression (.+-.error) measured for each of
           the tested oligomers.
           . . . oligomer. The position of the phosphorothicate core and/or the base composition of the phosphorothicate core has a large effect on
DETD
              ***antisense*** activity, as seen by comparing 3637-1, 3638-1, 3262-5
           and 3636-1. A more central position within the chimera is most active,.
                        (denoted by an "x" above the sequences shown above) within the
DETD
          RNaseH phosphorothioate core sequence of the chimeric oligomers eliminates ***antisense*** activity in this eukaryotic cell culture assay, as seen by comparing 3639-1 and 3640-1 with 3262-5. In a separate experiment.
          D. Demonstration of Activity of ***Antisense*** Chir
Targeted to HPV-11 E7 in Cell Free Translation Extracts
E. Demonstration of Activity of ***Antisense*** Olio
Cell-Free RNAseH Cleavage Assay
DETD
                                                                                          Chimeric Oligomers
DETD
                                                                                          Oligomers in a
DETD
           F. Demonstration of Activity of
                                                               ***Antisense***
                                                                                          Oligomers in
          Transiently Transfected COS-7 Cells
          Representative experiments were performed as follows. E7 expression plasmid pcDNA11E7 (5 .mu.g/ml) and different amounts of ***antisense*** oligonucleotide were transfected into COS-7 cells in
DETD
           the presence of Transfectam.TM. (Promega). Cells were incubated with
          DETD
                                                                                          Oligomers Targeted
           to E2 in Cell-Free Translation Extracts
```

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=> d history
            (FILE 'HOME' ENTERED AT 19:01:02 ON 21 JAN 2003)
           FILE 'MEDLINE, CAPLUS, LIFESCI, EMBASE, USPATFULL, BIOSIS' ENTERED AT 19:01:24 ON 21 JAN 2003
                         4478 S P53 AND ANTISENSE
L1
L2
                             78 S L1 AND SPLICE ACCEPTOR SITE
L3
                             78 DUP REM L2 (0 DUPLICATES REMOVED)
                               2 S L3 AND MORPHOLINO
=> s l1 and morpholino
                        171 L1 AND MORPHOLINO
=> dup rem 15
PROCESSING COMPLETED FOR L5
                          157 DUP REM L5 (14 DUPLICATES REMOVED)
=> s 16 and py<2001
       3 FILES SEARCHED...
                          37 L6 AND PY<2001
=> d 17 ibib abs tot
          ANSWER 1 OF 37
                                                    MEDLINE
                                            2000153646
ACCESSION NUMBER:
                                                                            MEDLINE
                                            20153646 PubMed ID: 10688605
c-Myc ***antisense*** limi
DOCUMENT NUMBER:
TITLE:
                                                                                                    limits rat liver regeneration and
                                            indicates role for c-Myc in regulating cytochrome P-450 3A
                                            activity.
                                           Arora V; Knapp D C; Smith B L; Statdfield M L; Stein D A; Reddy M T; Weller D D; Iversen P L AVI BioPharma, Corvallis, Oregon, USA. GM54871 (NIGMS)
AUTHOR:
CORPORATE SOURCE:
CONTRACT NUMBER:
                                            JOURNAL OF PHARMACOLOGY AND EXPERIMENTAL THERAPEUTICS, ***(2000 Mar)*** 292 (3) 921-8.
Journal code: 0376362. ISSN: 0022-3565.
SOURCE:
                                           United States
PUB. COUNTRY:
DOCUMENT TYPE:
                                            Journal; Article; (JOURNAL ARTICLE)
LANGUAGE:
                                           English
FILE SEGMENT:
                                            Priority Journals
ENTRY MONTH:
                                           200003
ENTRY DATE:
                                            Entered STN: 20000330
                                            Last Updated on STN: 20000330
          Expression of c-myc protein is associated with cell proliferation. The present study uses ***antisense*** oligomers to inhibit c-myc
AB
           expression in the regenerating rat liver after 70% partial hepatectomy
          (PH). ***Antisense*** phosphorodiamidate ***morpholino*** oligomers (novel DNA analogs) were administered i.p. immediately after surgery to block expression of c-myc within the first 24 h after PH. A
         surgery to block expression of c-myc within the first 24 h after PH. A 20-mer PMO complimentary to the c-myc mRNA at the translation start site was an effective sequence (AVI-4126, 5'-ACGTTGAGGGGCATCGTCGC-3'). A single i.p. dose of 0.5 mg/kg AVI-4126 caused reduction of the regenerating liver c-myc protein in a sequence-specific and dose-dependent manner. Inhibition of c-myc expression resulted in reduction of proliferating cell nuclear antigen and arrested cells in the G(0)/G(1) phase of the cell cycle. The ratio of G(2):G(0) cell populations in the regenerating liver 24 h after PH dropped from 29.1 in saline vehicle-treated rats to 18.0 in rats treated with 2.5 mg/kg AVI-4126. The expression of cell cycle checkpoint protein ***p53*** was inhibited with increasing doses of AVI-4126, but expression of p21(waf-1) was unaffected. The activity of cytochrome P-450 3A2 (CYP3A2) was evaluated by immunoblot analysis and erythromycin N-demethylation. AVI-4126 did not alter CYP3A activity in nonhepatectamized animals but showed a dose-dependent decrease in PH rats. We conclude that AVI-4126, ***antisense*** oligomer to c-myc, can reduce cell proliferation in the regenerating rat liver. Furthermore,
```

ANSWER 2 OF 37 CAPLUS COPYRIGHT 2003 ACS SSION NUMBER: 2000:291227 CAPLUS ACCESSION NUMBER: DOCUMENT NUMBER: 132:318574

TITLE: ***p53*** -targeted ***antisense***

We conclude that AVI-4126, ***antisense*** oligomer to c-myc, can reduce cell proliferation in the regenerating rat liver. Furthermore, inhibition of c-myc may indirectly influence the expression of CYP3A.

cancer and hypoxia-induced disorders

INVENTOR(S):

PATENT ASSIGNEE(S): SOURCE:

Iversen, Patrick L. Avi Biopharma, Inc., USA PCT Int. Appl., 27 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

APPLICATION NO. PATENT NO. KIND DATE DATE 20000504 wo 1999-us24758 19991022 <-wo 2000024885 Α2

wo 2000024885 Α3 20000720

W: AU, CA, JP, KR RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL,

PT, SE EP 1999-971033 EP 1124950 Α2 20010822 19991022

R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,

IE, FI

us 6365577 20020402 us 1999-426804 в1 19991022

PRIORITY APPLN. INFO.:

US 1998-105695P P 19981026 WO 1999-US24758 W 19991022

Antisense oligonucleotides useful for treating a disease state ***p53*** induction, such as proliferative cell characterized by disorders, e.g. cancer, or a hypoxic state induced by an ischemic attack, such as stroke, are described. The ***antisense*** agents are preferably of the class known as "steric blocker" type oligonucleotides, including ***morpholino*** oligonucleotides, peptide nucleic acids, 2' O-allyl or 2'-O-alkyl modified oligonucleotides, or N3' <<rwap P5' phosphoramidate oligonucleotides. Thus, in partially hepatectomized rats, ***morpholino*** - or C5-propyne cytosine-contg. ***p53***
antisense oligonucleotides enhanced wt. gain in the regenerating livers.

ANSWER 3 OF 37 USPATFULL

ACCESSION NUMBER:

2000:157559 USPATFULL

TITLE:

Modified oligonucleotides, their preparation and their

INVENTOR(S):

Seela, Frank, Osnabruck, Germany, Federal Republic of

<--

PATENT ASSIGNEE(S):

Thomas, Horst, Hasbergen, Germany, Federal Republic of Aventis Pharma Deutschland GmbH, Frankfurt am Main, Germany, Federal Republic of (non-U.S. corporation)

KIND NUMBER DATE

PATENT INFORMATION: APPLICATION INFO.:

US 6150510 20001121

RELATED APPLN. INFO.:

US 1998-144112 19980831 (9)

Continuation of Ser. No. US 1995-554164, filed on 6 Nov 1995, now patented, Pat. No. US 5844106

DOCUMENT TYPE: Utility FILE SEGMENT: Granted Riley, Jezia Foley & Lardner PRIMARY EXAMINER: LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:** 3592 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Modified oligonucleotides which possess at least one substituted 7-deazapurine base form more stable hybridization complexes with nucleic acids than unsubstituted analogs. They are useful as inhibitors of gene expression, as probes for detecting nucleic acids, as aids in molecular biology and as pharmaceuticals or diagnostic agents. Processes for preparing them are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 4 OF 37 USPATFULL

ACCESSION NUMBER:

2000:138328 USPATFULL

TITLE:

Inhibition of extracellular matrix synthesis by ***antisense*** compounds directed to nuclear

proto-oncogenes

INVENTOR(S):

Zalewski, Āndrew, Elkins Park, PA, United States

Shi, Yi, Cheltenham, PA, United States

PATENT ASSIGNEE(S):

Thomas Jefferson University, Philadelphia, PA, United

States (U.S. corporation)

NUMBER KIND DATE

US 6133242 20001017 PATENT INFORMATION: <--APPLICATION INFO.: us 1995-461366 19950605 (8)

Continuation-in-part of Ser. No. wo 1994-US11853, filed on 17 Oct 1994 which is a continuation-in-part of Ser. RELATED APPLN. INFO.:

No. US 1993-138637, filed on 15 Oct 1993, now abandoned

Utility DOCUMENT TYPE: FILE SEGMENT: Granted

Brusca, John S. PRIMARY EXAMINER: ASSISTANT EXAMINER: McGarry, Sean

Seidal, Gonda, Lavorgna & Monaco, PC LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 25 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 32 Drawing Figure(s); 23 Drawing Page(s)

2109 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A method and compounds are provided for inhibiting the synthesis of extracellular matrix proteins. Compounds of the invention comprise ***antisense*** oligonucleotides specific for nuclear proto-oncogenes. ΑB

antisense compounds of the invention are selected

from the group consisting of c-myc and c-myb and are locally administered. The invention finds use in the treatment of a variety of disorders, including sclerotic disorders and restenosis, associated with

the inappropriate synthesis of extracellular matrix proteins,

particularly collagen.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 5 OF 37 USPATFULL L7

ACCESSION NUMBER:

2000:124778 USPATFULL Antitumor ***antisense*** TITLE: sequences directed

against R1 and R2 components of ribonucleotide

reductase

INVENTOR(S): Wright, Jim A., Toronto, Canada Young, Aiping H., Toronto, Canada

PATENT ASSIGNEE(S): Genesense Technologies, Inc., Toronto, Canada (non-U.S.

corporation)

NUMBER KIND DATE

PATENT INFORMATION: us 6121000 20000919 <--

us 1999-249730 APPLICATION INFO.: 19990211 (9)

> NUMBER DATE

PRIORITY INFORMATION: US 1996-23040P 19960802 (60)

19970307 (60) US 1997-39959P Utility

DOCUMENT TYPE: FILE SEGMENT: Granted PRIMARY EXAMINER: Guzo, David ASSISTANT EXAMINER: Wang, Andrew

LEGAL REPRESENTATIVE: Burns, Doane, Swecker & Mathis, LLP

NUMBER OF CLAIMS: 11

EXEMPLARY CLAIM:

23 Drawing Figure(s); 23 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 4251

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Compounds and methods for modulating cell proliferation, preferably inhibiting the proliferation of tumor cells are described. Compounds

that may be used to modulate cell proliferation include
antisense oligonucleotides complementary to re

oligonucleotides complementary to regions of the

mammalian ribonucleotide reductase genes.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L7 ANSWER 6 OF 37 USPATFULL

ACCESSION NUMBER: 2000:109600 USPATFULL

Oligoribonucleotides and ribonucleases for cleaving RNA TITLE:

INVENTOR(S):

Crooke, Stanley T., Carlsbad, CA, United States
Isis Pharmaceuticals, Inc., Carlsbad, CA, United States PATENT ASSIGNEE(S):

(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 6107094 20000822

Continuation-in-part of Ser. No. US 1996-659440, filed RELATED APPLN. INFO.:

on 6 Jun 1996, now patented, Pat. No. US 5898031

Utility DOCUMENT TYPE: FILE SEGMENT: Granted

Elliott, George C. PRIMARY EXAMINER:

ASSISTANT EXAMINER: McGarry, Sean

LEGAL REPRESENTATIVE: Woodcock Washburn Kurtz Mackiewicz & Norris LLP

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 15 Drawing Figure(s); 10 Drawing Page(s)

LINE COUNT: 3806

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Oligomeric compounds including oligoribonucleotides and oligoribonucleosides are provided that have subsequences of 2'-pentoribofuranosyl nucleosides that activate dsRNase. The oligoribonucleotides and oligoribonucleosides can include substituent groups for increasing binding affinity to complementary nucleic acid strand as well as substituent groups for increasing nuclease resistance. The oligomeric compounds are useful for diagnostics and other research purposes, for modulating the expression of a protein in organisms, and for the diagnosis, detection and treatment of other conditions susceptible to oligonucleotide therapeutics. Also included in the invention are mammalian ribonucleases, i.e., enzymes that degrade RNA, and substrates for such ribonucleases. Such a ribonuclease is referred to herein as a dsRNase, wherein "ds" indicates the RNase's specificity for certain double-stranded RNA substrates. The artificial substrates for the dsRNases described herein are useful in preparing affinity matrices for purifying mammalian ribonuclease as well as non-degradative RNA-binding proteins.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 7 OF 37 USPATFULL

ACCESSION NUMBER: 2000:98562 USPATFULL

TITLE:

INVENTOR(S):

Circular DNA vectors for synthesis of RNA and DNA Kool, Eric T., Rochester, NY, United States University of Rochester, Rochester, NY, United States PATENT ASSIGNEE(S):

(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: APPLICATION INFO.:

us 6096880 20000801 US 1997-805631 19970226 (8)

RELATED APPLN. INFO.:

Continuation-in-part of Ser. No. US 1995-393439, filed on 23 Feb 1995, now patented, Pat. No. US 5714320 which is a continuation-part of Ser. No. US 1993-47860,

filed on 15 Apr 1993, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Elliot, George C. ASSISTANT EXAMINER: McGarry, Sean

Mueting, Raasch & Gebhardt, P.A. LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 8 Drawing Figure(s); 6 Drawing Page(s)

LINE COUNT: 3103

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides methods for synthesis, and therapeutic use of DNA and RNA oligonucleotides and analogs. RNA oligonucleotides arc synthesized using a small, circular DNA template which lacks an RNA polymerase promoter sequence. The RNA synthesis is performed by combining a circular single-stranded oligonucleotide template with an effective RNA polymerase and at least two types of ribonucleotide triphosphate to form an RNA oligonucleotide multimer comprising multiple copies of the desired RNA oligonucleotide sequence. Preferably, the RNA oligonucleotide multimer is cleaved to produce RNA oligonucleotides having well-defined ends. Preferred RNA oligonucleotide multimers contain ribozymes capable of both cis (autolytic) and trans cleavage.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 8 OF 37 USPATFULL L7

ACCESSION NUMBER: 2000:95115 USPATFULL

TITLE:

INVENTOR(S):

Cationic lipids
Lin, Kuei-Ying, Fremont, CA, United States
Mattuecci, Mark D., Burlingame, CA, United States

NUMBER KIND DATE us 6093816 20000725 PATENT INFORMATION: us 1996-672206 APPLICATION INFO.: 19960627 (8) DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Higel, Floyd D. woodcock Washburn Kurtz Mackiewicz & Norris LLP LEGAL REPRESENTATIVE: NUMBER OF CLAIMS: 34 **EXEMPLARY CLAIM:** 1 LINE COUNT: 2544 CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention is directed to new cationic lipids and intermediates in their synthesis that are useful for transfecting nucleic acids or peptides into prokaryotic or eukaryotic cells. The lipids comprise one or two substituted histidine residues, or similar compounds, linked to a lipophilic moiety. The lipids form a complex when mixed with polyanions such as nucleic acids or peptides. The complexes permit efficient transfer of polyanions into cells, usually without significant toxicity to the cells. CAS INDEXING IS AVAILABLE FOR THIS PATENT. ANSWER 9 OF 37 USPATFULL ١7 ACCESSION NUMBER: 2000:91721 USPATFULL Diagnostic method detecting loss of wild-type TITLE: ***p53*** INVENTOR(S): Vogelstein, Bert, Baltimore, MD, United States Baker, Suzanne J., Baltimore, MD, United States Fearon, Eric R., Baltimore, MD, United States Nigro, Janice M., Baltimore, MD, United States Johns Hopkins University, Baltimore, MD, United States PATENT ASSIGNEE(S): (U.S. corporation) NUMBER KTND DATE PATENT INFORMATION: US 6090566 20000718 <--US 1995-459676 19950602 (8) Division of Ser. No. US 1993-47041, filed on 22 Mar 1993, now patented, Pat. No. US 5527676 And a APPLICATION INFO.: RELATED APPLN. INFO.: continuation of Ser. No. US 1992-928661, filed on 17 Aug 1992, now abandoned And a continuation of Ser. No. US 1989-446584, filed on 6 Dec 1989, now abandoned And a continuation-in-part of Ser. No. US 1989-330566, filed on 29 Mar 1989, now abandoned DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Eyler, Yvonne Banner & Witcoff, Ltd. LEGAL REPRESENTATIVE: NUMBER OF CLAIMS: 13 EXEMPLARY CLAIM: NUMBER OF DRAWINGS: 9 Drawing Figure(s); 9 Drawing Page(s) LINE COUNT: 1011 CAS INDEXING IS AVAILABLE FOR THIS PATENT. Methods and kits are provided for assessing mutations and/or loss of the ***p53*** gene in human tumors. Both deletion mutations and point ***p53*** are observed in the same human tumor cells mutations in and these mutations are clonal within the cells of the tumor. Loss of wild-type ***p53*** genes is responsible for neoplastic progression. CAS INDEXING IS AVAILABLE FOR THIS PATENT. ١7 ANSWER 10 OF 37 USPATFULL ACCESSION NUMBER: 2000:77184 USPATFULL Highly sensitive multimeric nucleic acid probes TITLE: Kool, Eric T., Rochester, NY, United States University of Rochester, Rochester, NY, United States INVENTOR(S): PATENT ASSIGNEE(S): (U.S. corporation)

KIND

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DATE PATENT INFORMATION: us 6077668 us 1997-910632 20000620 (8) APPLICATION INFO.: 19970813

NUMBER

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1997-805631, filed Feb 1995, now patented, Pat. No. US 5714320, issued on 3 Feb 1998 which is a continuation-in-part of Ser. No. US 1993-47860, filed on 15 Apr 1993, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Brusca, John S. ASSISTANT EXAMINER:

McGarry, Sean Mueting, Raasch & Gebhardt, P.A. LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 8 Drawing Figure(s): 8 Drawing Page(s)

3477 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides detectably labeled RNA and DNA oligonucleotide multimers useful as diagnostic probes in medical, biological and chemical applications. A method for synthesizing DNA and RNA oligonucleotides, oligonucleotide multimers, and analogs, preferably those that are detectably labeled, is also provided. Oligonucleotide synthesis is performed by combining a circular single-stranded oligonucleotide template with an effective polymerase and at least two types of nucleotide triphosphate, without the addition of auxiliary proteins, to yield an oligonucleotide multimer comprising multiple copies of a repeated oligonucleotide sequence.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 11 OF 37 USPATFULL

2000:64945 USPATFULL ACCESSION NUMBER:

Modified oligonucleotides, their preparation and their TITLE:

Seela, Frank, Osnabruck, Germany, Federal Republic of Lampe, Sigrid, Berge/Hekese, Germany, Federal Republic INVENTOR(S):

PATENT ASSIGNEE(S): Hoechst Aktiengesellschaft, Frankfurt am Main, Germany,

Federal Republic of (non-U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: us 6066720 20000523 <--

APPLICATION INFO.: US 1998-94405 19980610 (9)

Division of Ser. No. US 1997-940196, filed on 29 Sep 1997, now patented, Pat. No. US 5789562 which is a continuation of Ser. No. US 1995-431777, filed on 1 May **RELATED APPLN. INFO.:**

1995, now abandoned

NUMBER DATE

PRIORITY INFORMATION: DE 1994-4415370 19940502

DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Riley, Jezia

LEGAL REPRESENTATIVE: Finnegan, Henderson, Farabow, Garrett & Dunner, L.L.P.

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 19 LINE COUNT: 2037

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The invention relates to novel modified oligonucleotides which contain AB at least one 8-azapurine base and form more stable hybridization complexes with nucleic acids; To a process for their preparation, and to their use as inhibitors of gene expression, as probes for detecting nucleic acids, as aids in molecular biology, and as a pharmaceutical or diagnostic agent.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 12 OF 37 USPATFULL L7

ACCESSION NUMBER:

2000:57750 USPATFULL Chimeric oligonucleoside compounds TITLE:

Arnold, Jr., Lyle J., Poway, CA, United States INVENTOR(S): Reynolds, Mark A., San Diego, CA, United States

Giachetti, Cristina, Solano Beach, CA, United States Lebedev, Alexandre V., San Diego, CA, United States Genta Incorporated, Lexington, MA, United States (U.S.

PATENT ASSIGNEE(S): corporation)

> NUMBER KIND DATE

PATENT INFORMATION: us 6060456 20000509

us 1997-960111 19971027 APPLICATION INFO.: (8) RELATED APPLN. INFO.:

Continuation of Ser. No. US 1995-481637, filed on 7 Jun 1995, now abandoned which is a continuation of Ser. No. US 1994-238177, filed on 4 May 1994, now abandoned which is a continuation of Ser. No. US 1994-233778,

filed on 26 Apr 1994, now abandoned which is a continuation of Ser. No. US 1993-154013, filed on 16 Nov 1993, now abandoned which is a continuation of Ser. No. US 1993-154014, filed on 16 Nov 1993, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Riley, Jezia

LEGAL REPRESENTATIVE: Knobbe, Martens, Olson & Bear LLP.

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 15 Drawing Figure(s); 12 Drawing Page(s)

LINE COUNT: 5081

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Chimeric oligonucleoside compounds, and methods of preparing and formulating the same, are disclosed. The compounds and compositions are useful in activating RNaseH-mediated cleavage of target ribonucleic acid

sequences, and in treating disease conditions relating to such

sequences.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 13 OF 37 USPATFULL

2000:41165 USPATFULL ACCESSION NUMBER:

TITLE: ***Antisense*** modulation of MDMX expression INVENTOR(S):

Monia, Brett P., La Costa, CA, United States Cowsert, Lex M., Carlsbad, CA, United States Isis Pharmaceuticals Inc., Carlsbad, CA, United States PATENT ASSIGNEE(S):

(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 6046320 20000404 <--APPLICATION INFO.: US 1999-289267 19990409 (9)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Elliott, George C.

ASSISTANT EXAMINER: Epps, Janet

Law Offices of Jane Massey Licata LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 14 **EXEMPLARY CLAIM:** 3298 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Antisense compounds, compositions and methods are provided for

modulating the expression of MDMX. The compositions comprise

antisense compounds, particularly ***antisense***
oligonucleotides, targeted to nucleic acids encoding MDMX. Methods of using these compounds for modulation of MDMX expression and for treatment of diseases associated with expression of MDMX are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 14 OF 37 USPATFULL

ACCESSION NUMBER: 2000:1692 USPATFULL

Sequence-directed DNA binding molecules compositions TITLE:

and methods

INVENTOR(S): Edwards, Cynthia A., Menlo Park, CA, United States

Cantor, Charles R., Boston, MA, United States
Andrews, Beth M., Maynard, MA, United States
Turin, Lisa M., Redwood City, CA, United States
Fry, Kirk E., Palo Alto, CA, United States

PATENT ASSIGNEE(S): Genelabs Technologies, Inc., Redwood, CA, United States

(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 6010849 20000104

APPLICATION INFO.: US 1995-482080 19950607

Division of Ser. No. US 1993-171389, filed on 20 Dec 1993, now patented, Pat. No. US 5578444 which is a continuation-in-part of Ser. No. US 1993-123936, filed RELATED APPLN. INFO.:

is a continuation-in-part of Ser. No. US 1992-996783 filed on 23 Dec 1992, now patented, Pat. No. US 5693463

which is a continuation-in-part of Ser. No. US 1991-723618, filed on 27 Jun 1991, now abandoned

DOCUMENT TYPE: Utility Granted FILE SEGMENT: PRIMARY EXAMINER: Degen, Nancy

ASSISTANT EXAMINER: Schwartzman, Robert

Fabin, Gary R.Dehlinger & Associates LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

48 Drawing Figure(s); 47 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 10022

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention defines a DNA:protein-binding assay useful for screening libraries of synthetic or biological compounds for their ability to bind DNA test sequences. The assay is versatile in that any number of test sequences can be tested by placing the test sequence adjacent to a defined protein binding screening sequence. Binding of molecules to these test sequence changes the binding characteristics of the protein molecule to its cognate binding sequence. When such a molecule binds the test sequence the equilibrium of the DNA:protein complexes is disturbed, generating changes in the concentration of free DNA probe. Numerous exemplary target test sequences (SEQ ID NO:1 to SEQ ID NO:600) are set forth. The assay of the present invention is also useful to characterize the preferred binding sequences of any selected DNA-binding molecule.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 15 OF 37 USPATFULL L7

ACCESSION NUMBER:

TITLE:

1999:159999 USPATFULL Antitumor ***antisense*** sequences directed

against ribonucleotide reductase

INVENTOR(S):

Wright, Jim A., 15 Bryn Mawr Road, Winnipeg, Manitoba.

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Canada R3T 3K8

Young, Aiping H., 717 Pacific Avenue, Winnipeg, Manitoba, Canada R3E 1G1

NUMBER KIND DATE

PATENT INFORMATION: US 5998383 19991207

US 1997-904901 APPLICATION INFO.: 19970801 (8)

> NUMBER DATE

US 1996-23040P US 1997-39959P 19960802 (60) 19970307 (60) PRIORITY INFORMATION:

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: LeGuyader, John L. ASSISTANT EXAMINER: Shibuya, Mark L. LEGAL REPRESENTATIVE: Burns, Doane, Swecker & Mathis, L.L.P.

NUMBER OF CLAIMS: 31 EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

10 Drawing Figure(s); 11 Drawing Page(s)

LINE COUNT: 4530

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

antisense A synthetic oligonucleotide comprising at least seven nucleotides or nucleotide analogues having a sequence complementary to the mRNA sequence of ribonucleotide reductase dimeric protein component R2 including SEQ ID Nos:1-102 is disclosed. A ***antisense*** oligonucleotide comprising at least seven nucleotides or nucleotide analogues having a sequence complementary to the mRNA sequence of ribonucleotide reductase dimeric protein component R1 including SEQ ID Nos:103-161 is also disclosed. The invention also discloses pharmaceutical compositions including the synthetic ***antisense***

oligonucleotides of the present invention and methods ***antisense*** oligonucleotides to modulation of using the

proliferative cells including neoplastic cells.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 16 OF 37 USPATFULL

ACCESSION NUMBER: 1999:159764 USPATFULL

Antisense TITLE: modulation of microtubuleINVENTOR(S): Bennett, C. Frank, Carlsbad, CA, United States

Ackermann, Elizabeth J., Solana Beach, CA, United

States

PATENT ASSIGNEE(S):

Isis Pharmaceuticals Inc., Carlsbad, CA, United States

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(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION:

us 5998148 us 1999-289368 19991207 19990408 (9)

APPLICATION INFO.: DOCUMENT TYPE:

Utility Granted

FILE SEGMENT: PRIMARY EXAMINER:

Elliot, George C.

ASSISTANT EXAMINER: LEGAL REPRESENTATIVE: Wang, Andrew

NUMBER OF CLAIMS:

Law Offices of Jane Massey Licata 14

EXEMPLARY CLAIM: LINE COUNT:

3094

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Antisense compounds, compositions and methods are provided for modulating the expression of microtubule-associated protein 4. The compositions comprise ***antisense*** compounds, particularly

antisense oligonucleotides, targeted to nucleic acids encoding
microtubule-associated protein 4. Methods of using these compounds for

modulation of microtubule-associated protein 4 expression and for treatment of diseases associated with expression of microtubule-

associated protein 4 are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 17 OF 37 USPATFULL L7

ACCESSION NUMBER:

1999:151195 USPATFULL

TITLE:

GATA-6 transcription factor: compositions and methods Walsh, Kenneth, Carlisle, MA, United States St. Elizabeth's Medical Center, Boston, MA, United

INVENTOR(S): PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: APPLICATION INFO.:

us 5990092 us 1997-927394 utility

19991123 19970827 (8)

DOCUMENT TYPE: FILE SEGMENT: PRIMARY EXAMINER:

Granted

ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: NUMBER OF CLAIMS:

Degen, Nancy Schwartzman, Robert Wolf, Greenfield & Sacks P.C. 21

EXEMPLARY CLAIM: LINE COUNT:

2449

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Methods and compositions for reducing or preventing the proliferation of vascular smooth muscle cells are provided. The method involves the step of administering an isolated GATA-6 molecule to a subject to prevent or reduce vascular smooth muscle cell proliferation. The isolated GATA-6 molecule can be a GATA-6 nucleic acid or a GATA-6 protein.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 18 OF 37 USPATFULL

ACCESSION NUMBER:

1999:96216 USPATFULL

TITLE:

Reagents and methods useful for detecting diseases of

the lung

INVENTOR(S):

Cohen, Maurice, Highland Park, IL, United States Friedman, Paula N., Deerfield, IL, United States Gordon, Julian, Lake Bluff, IL, United States Hodges, Steven C., Buffalo Grove, IL, United States Klass, Michael R., Libertyville, IL, United States Kratochvil, Jon D., Kenosha, WI, United States
Roberts-Rapp, Lisa, Gurnee, IL, United States
Russell, John C., Kenosha, WI, United States
Stroupe, Steven D., Libertyville, IL, United States

PATENT ASSIGNEE(S):

Abbott Laboratories, Abbott Park, IL, United States

(U.S. corporation)

NUMBER KIND DATE APPLICATION INFO.:

US 1997-964725 19971105 (8) Continuation-in-part of Ser. No. US 1996-744211, filed RELATED APPLN. INFO.:

on 5 Nov 1996, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Degren, Nancy

Wang, Andrew ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Becker, Cheryl L., Goller, Mimi C.

NUMBER OF CLAIMS: 21

EXEMPLARY CLAIM: NUMBER OF DRAWINGS: 6 Drawing Figure(s); 6 Drawing Page(s)

LINE COUNT: 3052

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A set of contiguous and partially overlapping RNA sequences and polypeptides encoded thereby, designated as LU103 and transcribed from lung tissue is described. A fully sequenced clone representing the longest continuous sequence of LU103 also disclosed. These sequences are useful for detecting, diagnosing, staging, monitoring, prognosticating, preventing or treating, or determining the predisposition of an individual to diseases and conditions of the lung

such as lung cancer.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 19 OF 37 USPATFULL

ACCESSION NUMBER: 1999:50839 USPATFULL

Oligoribonucleotides for cleaving RNA TITLE:

INVENTOR(S):

Crooke, Stanley T., Carlsbad, CA, United States ISIS Pharmaceuticals, Inc., Carlsbad, CA, United States PATENT ASSIGNEE(S):

(U.S. corporation)

NUMBER KIND DATE

US 5898031 US 1996-659440 PATENT INFORMATION: 19990427 <--APPLICATION INFO.: 19960606 (8)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: LeGuyader, John L.

LEGAL REPRESENTATIVE: Woodcock Washburn Kurtz Mackiewicz & Norris LLP

NUMBER OF CLAIMS: 66

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 12 Drawing Figure(s); 5 Drawing Page(s)

LINE COUNT: 3150

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Oligomeric compounds including oligoribonucleotides and oligoribonucleosides are provided that have subsequences of 2-pentoribofuranosyl nucleosides that activate dsRNase. The oligoribonucleotides and oligoribonucleosides can include substituent groups for increasing binding affinity to complementary nucleic acid strand as well as substituent groups for increasing nuclease resistance. The oligomeric compounds are useful for diagnostics and other research purposes, for modulating the expression of a protein in organisms, and for the diagnosis, detection and treatment of other conditions susceptible to oligonucleotide therapeutics.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 20 OF 37 USPATFULL

ACCESSION NUMBER: 1999:24783 USPATFULL

TITLE:

Therapeutic oligonucleotides targeting the human MDR1

and MRP genes

INVENTOR(S):

Smith, Larry J., Omaha, NE, United States
The Board of Regents of the University of Nebraska,
Lincoln, NE, United States (U.S. corporation) PATENT ASSIGNEE(S):

NUMBER KIND DATE PATENT INFORMATION: US 5874567 19990223

US 1997-927561 APPLICATION INFO.: 19970908

Continuation of Ser. No. US 1995-487141, filed on 7 Jun RELATED APPLN. INFO.:

1995, now patented, Pat. No. US 5683987 which is a continuation-in-part of Ser. No. US 1994-379180, filed

on 12 Jul 1994, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

LeGuyader, John L. PRIMARY EXAMINER:

LEGAL REPRESENTATIVE: Dann, Dorfman, Herrell and Skillman

NUMBER OF CLAIMS: 15

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 1 Drawing Figure(s); 1 Drawing Page(s)

LINE COUNT:

2080

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides novel compositions and methods useful in cancer therapy for inhibiting the multidrug resistance phenotype, which often thwarts long-term chemotherapeutic regimens. The novel compositions of matter comprise oligonucleotides targeted to the human

MDR1 and MRP genes, which inhibit expression of these genes, thereby rendering tumors and other forms of cancer more susceptible to the cytotoxic effects of chemotherapeutic agents. Oligonucleotides are also provided that inhibit the multidrug resistance phenotype by exerting an aptameric effect.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 21 OF 37 USPATFULL

1999:18912 USPATFULL ACCESSION NUMBER:

Method of determining DNA sequence preference of a TITLE:

DNA-binding molecule

INVENTOR(S):

Edwards, Cynthia A., Menlo Park, CA, United States
Cantor, Charles R., Boston, MA, United States
Andrews, Beth M., Maynard, MA, United States
Turin, Lisa M., Redwood City, CA, United States
Fry, Kirk E., Palo Alto, CA, United States
Genelabs Technologies, Inc., Redwood City, CA, United
States (U.S. Corporation)

PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 5869241 19990209

US 1995-475228 APPLICATION INFO.: 19950607 RELATED APPLN. INFO.:

Division of Ser. No. US 1993-171389, filed on 20 Dec 1993, now patented, Pat. No. US 5578444 which is a continuation-in-part of Ser. No. US 1993-123936, filed on 17 Sep 1993, now patented, Pat. No. US 5726014 which is a continuation-in-part of Ser. No. US 1992-996783, filed on 23 Dec 1992, now patented, Pat. No. US 5693463 which is a continuation-in-part of Ser. No. US 1991-723618, filed on 27 Jun 1991, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Zitomer, Stepanie W. Whisenant, Ethan ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Fabian, Gary R., Stratford, Carol A., Dehlinger, Peter

11 NUMBER OF CLAIMS: EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 72 Drawing Figure(s); 47 Drawing Page(s)

LINE COUNT: 9840

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention defines a DNA:protein-binding assay useful for screening libraries of synthetic or biological compounds for their ability to bind DNA test sequences. The assay is versatile in that any number of test sequences can be tested by placing the test sequence adjacent to a defined protein binding screening sequence. Binding of molecules to these test sequence changes the binding characteristics of the protein molecule to its cognate binding sequence. When such a molecule binds the test sequence the equilibrium of the DNA: protein complexes is disturbed, generating changes in the concentration of free DNA probe. Numerous exemplary target test sequences (SEQ ID NO:1 to SEQ ID NO:600) are set forth. The assay of the present invention is also useful to characterize the preferred binding sequences of any selected DNA-binding molecule.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 22 OF 37 USPATFULL

1999:4647 USPATFULL ACCESSION NUMBER:

Fas ligand compositions for treatment of proliferative TITLE:

disorders

INVENTOR(S):

Walsh, Kenneth, Carlisle, MA, United States St. Elizabeth's Medical Center, Boston, MA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE us 5858990 PATENT INFORMATION: 19990112 APPLICATION INFO.: US 1997-810453 19970304 (8) DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Elliott, George C. ASSISTANT EXAMINER: McGarry, Sean LEGAL REPRESENTATIVE: Wolf, Greenfield & Sacks, P.C. NUMBER OF CLAIMS: **EXEMPLARY CLAIM:** 3038 LINE COUNT: CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB A method for treating vascular injury, particularly vascular injury resulting from restenosis following angioplasty, and vascular remodeling is provided. The method involves administering to subjects in need of such treatment an effective amount of a Fas ligand molecule. CAS INDEXING IS AVAILABLE FOR THIS PATENT. ANSWER 23 OF 37 USPATFULL L7 ACCESSION NUMBER: 1998:151104 USPATFULL TITLE: Modified oligonucleotides, their preparation and their use Seela, Frank, Osnabrueck, Germany, Federal Republic of INVENTOR(S): Thomas, Horst, Hasbergen, Germany, Federal Republic of Hoechst Aktiengesellschaft, Germany, Federal Republic PATENT ASSIGNEE(S): of (non-U.S. corporation) NUMBER KIND DATE US 5844106 US 1995-554164 PATENT INFORMATION: 19981201 <--APPLICATION INFO.: 19951106 (8) DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Marschel, Ardin H. ASSISTANT EXAMINER: Riley, Jézia Foley & Lardner LEGAL REPRESENTATIVE: NUMBER OF CLAIMS: 17 **EXEMPLARY CLAIM:** 1 LINE COUNT: 2731 CAS INDEXING IS AVAILABLE FOR THIS PATENT. AB Modified oligonucleotides which possess at least one substituted 7-deazapurine base form more stable hybridization complexes with nucleic acids than unsubstituted analogs. They are useful as inhibitors of gene expression, as probes for detecting nucleic acids, as aids in molecular biology and as pharmaceuticals or diagnostic agents. Processes for preparing CAS INDEXING IS AVAILABLE FOR THIS PATENT. ANSWER 24 OF 37 USPATFULL ACCESSION NUMBER: 1998:150686 USPATFULL TITLE: Cleavage of nucleic acid acid using thermostable methoanococcus jannaschii FEN-1 endonucleases Kaiser, Michael W., Madison, WI, United States INVENTOR(S): Lyamichev, Victor I., Madison, WI, United States Lyamichev, Natasha, Madison, WI, United States PATENT ASSIGNEE(S): Third wave Technologies, Inc., Madison, WI, United States (U.S. corporation) NUMBER KIND DATE PATENT INFORMATION: US 5843669 19981201 <--APPLICATION INFO.: US 1996-757653 19961129 (8) RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1996-599491, filed on 24 Jan 1996 DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Jones, W. Gary

ASSISTANT EXAMINER: Fredman, Jeffrey LEGAL REPRESENTATIVE: Medlen & Carroll, LLP NUMBER OF CLAIMS: 26 **EXEMPLARY CLAIM:** NUMBER OF DRAWINGS: 161 Drawing Figure(s); 131 Drawing Page(s) LINE COUNT: 15189

AB The present invention relates to means for cleaving a nucleic acid cleavage structure in a site-specific manner. Structure-specific nucleases, including 5' nucleases, thermostable FEN-1 endonucleases and 3' exonucleases, are used to detect and identify target nucleic acids. Methods are provided which allow for the detection specific nucleic acid sequences; these methods permit the detection and identification of mutant and wild-type forms of genes (e.g., human genes) as well as permit the detection and identification of bacterial and viral pathogens in a sample.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 25 OF 37 USPATFULL

ACCESSION NUMBER: 1998:150671 USPATFULL

TITLE: INVENTOR(S): Rapid detection of mutations in the ***p53*** aene

Heisler, Laura M., Madison, WI, United States
Fors, Lance, Monrovia, CA, United States
Brow, Mary Ann D., Madison, WI, United States
Third Wave Technologies, Inc., Madison, WI, United
States (U.S. corporation)

PATENT ASSIGNEE(S):

DATE NUMBER KIND _____

PATENT INFORMATION: APPLICATION INFO.: RELATED APPLN. INFO.: US 5843654 19981201 US 1995-484956 19950607 (8)

Continuation-in-part of Ser. No. US 1995-402601, filed on 9 Mar 1995 which is a continuation-in-part of Ser. No. US 1994-337164, filed on 9 Nov 1994, now abandoned

which is a continuation-in-part of Ser. No. US 1994-254359, filed on 6 Jun 1994, now patented, Pat. No. US 5614402, issued on 25 Mar 1997 which is a continuation-in-part of Ser. No. US 1993-73384, file on 4 Jun 1993, now patented, Pat. No. US 5541311, issued on 30 Jul 1996 which is a continuation-in-part of Ser. No. US 1992-986330, filed on 7 Dec 1992, now patented, Pat. No. US 5422253, issued on 6 Jun 1995

DOCUMENT TYPE: Utility Granted

FILE SEGMENT: PRIMARY EXAMINER:

Patterson, Jr., Charles L. Medlen & Carroll, LLP

LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 25 **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 118 Drawing Figure(s); 79 Drawing Page(s)

LINE COUNT: 11066

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to means for cleaving a nucleic acid cleavage structure in a site-specific manner. Enzymes, including 5' nucleases and 3' exonucleases, are used to screen for known and unknown mutations, including single base changes, in the human ***p53*** mutations, including single base changes, in the human ***p53*** gene. Methods are provided which allow for the identification of genetic

p53 mutations in the human gene in a sample.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 26 OF 37 USPATFULL L7

ACCESSION NUMBER:

1998:92178 USPATFULL

TITLE:

Nucleotide monomers containing 8-azapurin bases or a derivative thereof, their preparation and their use in making modified olignonucleotides

INVENTOR(S):

Seela, Frank, Osnabruck, Germany, Federal Republic of Lampe, Sigrid, Berge/Hekese, Germany, Federal Republic

PATENT ASSIGNEE(S):

Hoechst Aktiengesellschaft, Frankfurt am Main, Germany,

<--

Federal Republic of (non-U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION:

us 5789562 19980804

APPLICATION INFO.:

us 1997-940196 19970929 (8)

RELATED APPLN. INFO.:

Continuation of Ser. No. US 1995-431777, filed on 1 May

1995, now abandoned

NUMBER DATE

PRIORITY INFORMATION:

DE 1994-4415370 19940502

DOCUMENT TYPE:

Utility

PRIMARY EXAMINER: Marschel, Ardin н.

ASSISTANT EXAMINER: Rilev, Jezia

LEGAL REPRESENTATIVE: Finnegan, Henderson, Farabow, Garrett & Dunner, L.L.P.

NUMBER OF CLAIMS: EXEMPLARY CLAIM: LINE COUNT: 1868

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The invention relates to novel modified oligonucleotides which contain at least one 8-azapurine base and form more stable hybridization complexes with nucleic acids; To a process for their preparation, and to their use as inhibitors of gene expression, as probes for detecting nucleic acids, as aids in molecular biology, and as a pharmaceutical or diagnostic agent.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 27 OF 37 USPATFULL

ACCESSION NUMBER: 1998:68824 USPATFULL

Targeted nucleic acid delivery into liver cells TITLE:

INVENTOR(S): Kuo, M. Tien, Houston, TX, United States Ding, Zhi-Ming, Houston, TX, United States

Board of Regents , The University of Texas System, PATENT ASSIGNEE(S):

Austin, TX, United States (U.S. corporation)

NUMBER KIND DATE

US 5766899 US 1995-395602 19980616 PATENT INFORMATION: <--19950227 (8)

APPLICATION INFO.: Utility DOCUMENT TYPE: FILE SEGMENT: Granted

Robinson, Douglas W. PRIMARY EXAMINER: ASSISTANT EXAMINER:

Nelson, Amy J. Arnold, White & Durkee LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 29 EXEMPLARY CLAIM: 26

NUMBER OF DRAWINGS: 20 Drawing Figure(s); 15 Drawing Page(s)

LINE COUNT: 1842

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Disclosed is a receptor-mediated complex that selectively delivers nucleic acid into hepatocytes. Circumsporozoite (CS) protein is the targeting ligand that recognizes a receptor expressed on the liver cell surface. The CS ligand is complexed with a polylysine component that can bind nucleic acid. The level of gene expression is greatly enhanced when the complex is cotransfected with adenovirus. Using the present invention, a reporter gene was successfully transferred into a number of different cell lines that express high levels of receptor. The ability to introduce nucleic acid into specific mammalian cells is an important therapy for numerous diseases such as cancer, malaria and hepatitis.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 28 OF 37 USPATFULL

ACCESSION NUMBER: 1998:44877 USPATFULL

TITLE: Sequence-directed DNA-binding molecules compositions

INVENTOR(S):

Edwards, Cynthia A., Menlo Park, CA, United States Fry, Kirk E., Palo Alto, CA, United States Cantor, Charles R., Boston, MA, United States Andrews, Beth M., Maynard, MA, United States Genelabs Technologies, Inc., Redwood City, CA, United

PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 5744131 19980428 <--APPLICATION INFO.: US 1995-476876 19950607 (8)

Division of Ser. No. US 1992-996783, filed on 23 Dec RELATED APPLN. INFO.: 1992 which is a continuation-in-part of Ser. No. US 1991-723618, filed on 27 Jun 1991, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Zitomer, Stephanie W.

3

ASSISTANT EXAMINER: Atzel, Amy

LEGAL REPRESENTATIVE: Fabian, Gary R., Stratford, Carol A., Dehlinger, Peter

NUMBER OF CLAIMS:

NUMBER OF DRAWINGS: 48 Drawing Figure(s); 33 Drawing Page(s)

LINE COUNT: 5113

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention defines an assay useful for screening libraries of synthetic or biological compounds for their ability to bind specific DNA test sequences. The assay is also useful for determining the sequence specificity and relative DNA-binding affinity of DNA-binding molecules for any particular DNA sequence. Also described herein are potential applications of the assay, including: 1) the detection of lead compounds or new drugs via the mass screening of libraries of synthetic or biological compounds (i.e., fermentation broths); 2) the design of sequence-specific DNA-binding drugs comprised of homo- or hetero-meric subunits of molecules for which the sequence specificity was determined using the assay; and 3) the use of molecules for which sequence specificity was determined using the assay as covalently attached moieties to aid in the binding of nucleic acid or other macromolecular polymers to nucleic acid sequences.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 29 OF 37 USPATFULL

ACCESSION NUMBER: 1998:39383 USPATFULL

Sequence-directed DNA-binding molecules compositions TITLE:

and methods

INVENTOR(S):

Edwards, Cynthia A., Menlo Park, CA, United States Fry, Kirk E., Palo Alto, CA, United States Cantor, Charles R., Boston, MA, United States Andrews, Beth M., Maynard, MA, United States

Genelabs Technologies, Inc., Redwood City, CA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 5738990 19980414 <--APPLICATION INFO.: US 1995-475221 19950607

Division of Ser. No. US 1992-996783, filed on 23 Dec 1992 which is a continuation-in-part of Ser. No. US 1991-723618, filed on 27 Jun 1991, now abandoned RELATED APPLN. INFO.:

DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Guzo, David ASSISTANT EXAMINER: Brusca, John S.

LEGAL REPRESENTATIVE: Fabian, Gary R., Stratford, Carol A., Dehlinger, Peter

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 48 Drawing Figure(s); 33 Drawing Page(s)

5040 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention defines an assay useful for screening libraries of synthetic or biological compounds for their ability to bind specific DNA test sequences. The assay is also useful for determining the sequence specificity and relative DNA-binding affinity of DNA-binding molecules for any particular DNA sequence. Also described herein are potential applications of the assay, including: 1) the detection of lead compounds or new drugs via the mass screening of libraries of synthetic or biological compounds (i.e., fermentation broths); 2) the design of sequence-specific DNA-binding drugs comprised of homo- or hetero-meric subunits of molecules for which the sequence specificity was determined using the assay; and 3) the use of molecules for which sequence specificity was determined using the assay as covalently attached mojeties to aid in the binding of nucleic acid or other macromolecular polymers to nucleic acid sequences.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 30 OF 37 USPATFULL

ACCESSION NUMBER: 1998:31048 USPATFULL

TITLE: Method of use of radicical for treatment of

immunopathological disorders

INVENTOR(S): Feng, Lili, San Diego, CA, United States

Hwang, Daniel, Baton Rouge, LA, United States The Scripps Research Institute, La Jolla, CA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

Board of Supervisors of Louisiana State University and Agricultural and Mechanical College, Baton Rouge, LA,

NUMBER KIND DATE US 5731343 US 1995-394148 PATENT INFORMATION: 19980324 APPLICATION INFO.: 19950224 (8)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

Cintins, Marianne M. Jarvis, William R. A. PRIMARY EXAMINER: ASSISTANT EXAMINER: Fish & Richardson, P.C. LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 43 Drawing Figure(s); 24 Drawing Page(s)

1540 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides a method of treating an immunopathological disorder having an etiology associated with production of a proinflammatory agent, by administering a compound of the formula: ##STR1## where R1 and R2 are independently H or --COR3; R3 is H, 1-50C alkyl, 1-20C alkoxy, 2-30C alkenyl, 2-30C alkenyloxy, 2-10 alkynyl, 6-14C aryl or aryloxy, a 5-6 membered heterocycle (containing 1-3 N, O and/or S heteroatoms and optionally fused to an aryl group), 3-8C cycloalkyl (optionally fused to aryl) or 5-8C cycloalkenyl; and R4 is a halogen. Examples of such proinflammatory agents include interlegistic 1. (7) interlegistic 1. (7) interlegistic 1. interleukin-1 (IL-1), interleukin-6 (IL-6), interferon-.gamma. (IFN-.gamma.), tumor necrosis factor-.alpha. (TNF-.alpha.), granulocyte macrophage-colony stimulating factor (GM-CSF), the growth related gene KC, cyclooxygenase-1 (COX-1), cyclooxygenase-2 (COX-2), macrophage chemotactic protein (MCP), inducible nitric oxide synthetase (iNOS), macrophage inflammatory protein (MIR) +iccur factor (TT) macrophage inflammatory protein (MIP), tissue factor (TF), phosphotyrosine phosphatase (PTPase), and endotoxin.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 31 OF 37 USPATFULL

ACCESSION NUMBER: 1998:25075 USPATFULL

TITLE: Screening assay for the detection of DNA-binding

molecules

INVENTOR(S):

Edwards, Cynthia A., Menlo Park, CA, United States Cantor, Charles R., Boston, MA, United States Andrews, Beth M., Watertown, MA, United States Turin, Lisa M., Berkeley, CA, United States

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Genelabs Technologies, Inc., Redwood City, CA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

US 5726014 PATENT INFORMATION: 19980310 19930917 APPLICATION INFO.: us 1993-123936 (8)

RELATED APPLN. INFO.:

Continuation-in-part of Ser. No. US 1992-996783, filed on 23 Dec 1992 which is a continuation-in-part of Ser. No. US 1991-723618, filed on 27 Jun 1991, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Jones, W. Gary Atzel, Amy ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Fabian, Gary R., Stratford, Carol A., Dehlinger, Peter

19 NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 72 Drawing Figure(s); 47 Drawing Page(s)

LINE COUNT: 5659

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention defines a DNA: protein-binding assay useful for screening libraries of synthetic or biological compounds for their ability to bind DNA test sequences. The assay is versatile in that any number of test sequences can be tested by placing the test sequence adjacent to a defined protein binding screening sequence. Binding of molecules to these test sequence changes the binding characteristics of the protein molecule to its cognate binding sequence. When such a molecule binds the test sequence the equilibrium of the DNA:protein complexes is disturbed, generating changes in the concentration of free DNA probe. Numerous exemplary target test sequences (SEQ ID NO:1 to SEQ ID NO:600) are set forth. The assay of the present invention is also useful to characterize the preferred binding sequences of any selected DNA-binding molecule.

ANSWER 32 OF 37 USPATFULL

ACCESSION NUMBER: 1998:14634 USPATFULL

TITLE: Method of constructing sequence-specific DNA-binding

molecules

INVENTOR(S):

Edwards, Cynthia A., Menlo Park, CA, United States Fry, Kirk E., Palo Alto, CA, United States Cantor, Charles R., Boston, MA, United States

Andrews, Beth M., Watertown, MA, United States Genelabs Technologies, Inc., Redwood City, CA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 5716780 19980210 <--

US 1995-484499 19950607 APPLICATION INFO.: RELATED APPLN. INFO.: Division of Ser. No. US 1992-996783, filed on 23 Dec

1992 which is a continuation-in-part of Ser. No. US 1991-723618, filed on 27 Jun 1991, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

Jones, W. Gary Atzel, Amy PRIMARY EXAMINER: ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Fabian, Gary R., Stratford, Carol A., Dehlinger, Peter

9 NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 48 Drawing Figure(s); 33 Drawing Page(s)

LINE COUNT: 4929

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention defines an assay useful for screening libraries of synthetic or biological compounds for their ability to bind specific DNA test sequences. The assay is also useful for determining the sequence specificity and relative DNA-binding affinity of DNA-binding molecules for any particular DNA sequence. Also described herein are potential applications of the assay, including: 1) the detection of lead compounds or new drugs via the mass screening of libraries of synthetic or biological compounds (i.e., fermentation broths); 2) the design of sequence-specific DNA-binding drugs comprised of homo- or hetero-meric subunits of molecules for which the sequence specificity was determined using the assay; and 3) the use of molecules for which sequence specificity was determined using the assay as covalently attached mojeties to aid in the binding of nucleic acid or other macromolecular polymers to nucleic acid sequences.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L7 ANSWER 33 OF 37 USPATFULL

ACCESSION NUMBER: 1998:11870 USPATFULL

Rolling circle synthesis of oligonucleotides and TITLE:

amplification of select randomized circular

oligonucleotides

INVENTOR(S):

Kool, Eric T., Rochester, NY, United States University of Rochester, Rochester, NY, United States PATENT ASSIGNEE(S):

(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 5714320 19980203

us 1995-393439 19950223 APPLICATION INFO.: (8)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1993-47860, filed

on 15 Apr 1993, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted Jones, W. Gary Rees, Dianne PRIMARY EXAMINER: ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Mueting, Raasch, Gebhardt & Schwappach, P.A.

NUMBER OF CLAIMS: 47 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 2 Drawing Figure(s); 2 Drawing Page(s)

LINE COUNT: 2583

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides methods for synthesis, selection, and AB amplification of DNA and RNA oligonucleotides and analogs. The method for synthesizing an oligonucleotide involves: providing an effective

at least one copy of the desired oligonucleotide sequence linked to a cleavage site; providing an effective amount of an isolated oligonucleotide primer; annealing the primer to the circular template to form a primed circular template; and combining the primed circular template with an effective amount of at least two types of nucleotide triphosphates and an effective amount of a polymerase enzyme to form a nucleotide multimer complementary to the circular oligonucleotide template, wherein the nucleotide multimer comprises multiple copies of the oligonucleotide sequence joined end to end. Preferably, the nucleotide multimer is cleaved to produce oligonucleotides having well-defined ends.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 34 OF 37 USPATFULL

ACCESSION NUMBER: 97:112300 USPATFULL

TITLE:

Method of ordering sequence binding preferences of a DNA-binding molecule

INVENTOR(S):

Edwards, Cynthia A., Menlo Park, CA, United States
Fry, Kirk E., Palo Alto, CA, United States
Cantor, Charles R., Boston, MA, United States
Andrews, Beth M., Maynard, MA, United States(4)
Genelabs Technologies, Inc., Redwood City, CA, United
States (U.S. corporation)

PATENT ASSIGNEE(S):

NUMBER DATE KIND

PATENT INFORMATION: us 5693463 19971202 <--

us 1992-996783 19921223 (7) APPLICATION INFO.:

DISCLAIMER DATE: 20110426

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1991-723618, filed

on 27 Jun 1991, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

Zitomer, Stephanie W. PRIMARY EXAMINER:

ASSISTANT EXAMINER: Atzel, Amy

LEGAL REPRESENTATIVE: Fabian, Gary R., Stratford, Carol A., Dehlinger, Peter

NUMBER OF CLAIMS: EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

48 Drawing Figure(s); 33 Drawing Page(s)

LINE COUNT: 4908

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention defines an assay useful for screening libraries of synthetic or biological compounds for their ability to bind specific DNA test sequences. The assay is also useful for determining the sequence specificity and relative DNA-binding affinity of DNA-binding molecules for any particular DNA sequence. Also described herein are potential applications of the assay, including: 1) the detection of lead compounds or new drugs via the mass screening of libraries of synthetic or biological compounds (i.e., fermentation broths); 2) the design of sequence-specific DNA-binding drugs comprised of homo- or hetero-meric subunits of malacular for which the comprise of provided the comprised control of the comprised of the comprised the comprised control of the control of the comprised control of the control o subunits of molecules for which the sequence specificity was determined using the assay; and 3) the use of molecules for which sequence specificity was determined using the assay as covalently attached molecules to aid in the binding of nucleic acid or other macromolecular

polymers to nucleic acid sequences.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L7 ANSWER 35 OF 37 USPATFULL

97:101743 USPATFULL ACCESSION NUMBER:

TITLE:

Therapeutic oligonucleotides targeting the human MDR1

<--

and MRP genes

INVENTOR(S): PATENT ASSIGNEE(S):

Smith, Larry J., Omaha, NE, United States The Board of Regents of the University of Nebraska, Lincoln, NE, United States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: us 5683987 19971104

US 1995-487141 19950607 (8) APPLICATION INFO.:

Continuation-in-part of Ser. No. US 1994-379180, filed RELATED APPLN. INFO.:

on 12 Jul 1994, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted ASSISTANT EXAMINER: Wang, Andrew

Dann, Dorfman, Herrell and Skillman LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 32 EXEMPLARY CLAIM:

2 Drawing Figure(s); 1 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 2111

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides novel compositions and methods useful in

cancer therapy for inhibiting the multidrug resistance phenotype, which often thwarts long-term chemotherapeutic regimens. The novel compositions of matter comprise oligonucleotides targeted to the human MDR1 and MRP genes, which inhibit expression of these genes, thereby rendering tumors and other forms of cancer more susceptible to the cytotoxic effects of chemotherapeutic agents. Oligonucleotides are also provided that inhibit the multidrug resistance phenotype by exerting an aptameric effect.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 36 OF 37 USPATFULL L7

ACCESSION NUMBER: 97:17918 USPATFULL

TITLE:

Compositions and methods for enhanced drug delivery

INVENTOR(S):

Hale, Ron L., Woodside, CA, United States
Lu, Amy, Los Altos, CA, United States
Solas, Dennis, San Francisco, CA, United States
Selick, Harold E., Belmont, CA, United States Oldenburg, Kevin R., Fremont, CA, United States Zaffaroni, Alejandro C., Atherton, CA, United States Affymax Technologies N.V., Middlesex, England (non-U.S.

PATENT ASSIGNEE(S):

corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 5607691 19970304 <--US 1995-449188 APPLICATION INFO.: 19950524 (8)

RELATED APPLN. INFO.: Continuation of Ser. No. US 1993-164293, filed on 9 Dec

1993, now abandoned which is a continuation-in-part of Ser. No. US 1993-77296, filed on 14 Jun 1993, now abandoned which is a continuation-in-part of Ser. No. US 1992-898219, filed on 12 Jun 1992, now abandoned And a continuation-in-part of Ser. No. US 1993-9463, filed

on 27 Jan 1993, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Levy, Neil S. LEGAL REPRESENTATIVE: Stevens, Lauren L.

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 5349 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to methods of delivering pharmaceutical agents across membranes, including the skin layer or mucosal membranes of a patient. A pharmaceutical agent is covalently bonded to a chemical modifier, via a physiologically cleavable bond, such that the membrane transport and delivery of the agent is enhanced.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 37 OF 37 USPATFULL

ACCESSION NUMBER: 96:108816 USPATFULL

Sequence-directed DNA-binding molecules compositions TITLE:

and methods

INVENTOR(S):

Edwards, Cynthia A., Menlo Park, CA, United States Cantor, Charles R., Boston, MA, United States Andrews, Beth M., Maynard, MA, United States Turin, Lisa M., Redwood City, CA, United States

Fry, Kirk E., Palo Alto, CA, United States

PATENT ASSIGNEE(S): Genelabs Technologies, Inc., Redwood City, CA, United

States (U.S. corporation)

NUMBER KIND DATE

US 5578444 US 1993-171389 PATENT INFORMATION: 19961126

(8) APPLICATION INFO.: 19931220 Continuation-in-part of Ser. No. US 1993-123936, filed RELATED APPLN. INFO.:

on 17 Sep 1993 which is a continuation-in-part of Ser.

continuation-in-part of Ser. No. US 1991-723618, filed

on 27 Jun 1991, now abandoned

DOCUMENT TYPE: Utility Granted FILE SEGMENT:

Zitomer, Stephanie W. PRIMARY EXAMINER:

ASSISTANT EXAMINER: Atzel, Amy

Fabian, Gary R., Brookes, Allen A., Stratford, Carol A. LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS:

EXEMPLARY CLAIM:

71 Drawing Figure(s); 48 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 5845

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention defines a DNA:protein-binding assay useful for screening libraries of synthetic or biological compounds for their ability to bind DNA test sequences. The assay is versatile in that any number of test sequences can be tested by plaining the test sequence. adjacent to a defined protein binding screening sequence. Binding of molecules to these test sequence changes the binding characteristics of the protein molecule to its cognate binding sequence. When such a molecule binds the test sequence the equilibrium of the DNA:protein complexes is disturbed, generating changes in the concentration of free DNA probe. Numerous exemplary target test sequences (SEQ ID NO:1 to SEQ ID NO:600) are set forth. The assay of the present invention is also useful to characterize the preferred binding sequences of any selected DNA-binding molecule.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

=> d history

L1

(FILE 'HOME' ENTERED AT 19:01:02 ON 21 JAN 2003)

FILE 'MEDLINE, CAPLUS, LIFESCI, EMBASE, USPATFULL, BIOSIS' ENTERED AT 19:01:24 ON 21 JAN 2003

4478 S P53 AND ANTISENSE

L2 78 S L1 AND SPLICE ACCEPTOR SITE

78 DUP REM L2 (0 DUPLICATES REMOVED)

L3 L4 2 S L3 AND MORPHOLINO

171 S L1 AND MORPHOLINO L5

L6 157 DUP REM L5 (14 DUPLICATES REMOVED)

37 S L6 AND PY<2001

=> s p53 (p) antisense

2189 P53 (P) ANTISENSE

=> s 18 and morpholino

66 L8 AND MORPHOLINO

=> s 19 not 17

L10 54 L9 NOT L7

=> dup rem 110

PROCESSING COMPLETED FOR L10

41 DUP REM L10 (13 DUPLICATES REMOVED)

=> d 111 ibib abs tot

L11 ANSWER 1 OF 41 USPATFULL

2003:17392 USPATFULL ACCESSION NUMBER:

HS2STs as modifiers of the p53 pathway and methods of TITLE:

use

INVENTOR(S): Friedman, Lori, San Francisco, CA, UNITED STATES

Plowman, Gregory D., San Carlos, CA, UNITED STATES Belvin, Marcia, Albany, CA, UNITED STATES

Francis-Lang, Helen, San Francisco, CA, UNITED STATES Li, Danxi, San Francisco, CA, UNITED STATES Funke, Roel P., South San Francisco, CA, UNITED STATES

NUMBER KIND DATE PATENT INFORMATION: US 2003013144 Α1 20030116 20020603 (10) APPLICATION INFO.: US 2002-161398 Α1

> NUMBER DATE

US 2001-328605P 20011010 (60) US 2002-357253P 20020215 (60)

DOCUMENT TYPE: Utility

APPLICATION FILE SEGMENT:

JAN P. BRUNELLE, EXELIXIS, INC., 170 HARBOR WAY, P.O. BOX 511, SOUTH SAN FRANCISCO, CA, 94083-0511 LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 25 **EXEMPLARY CLAIM:** 2023 LINE COUNT:

Human HS2ST genes are identified as modulators of the p53 pathway, and AB thus are therapeutic targets for disorders associated with defective p53

function. Methods for identifying modulators of p53, comprising

screening for agents that modulate the activity of HS2ST are provided.

L11 ANSWER 2 OF 41 USPATFULL **DUPLICATE 1**

ACCESSION NUMBER:

2002:69973 USPATFULL ***p53*** ***an1 ***antisense*** agent and method TITLE: Iversen, Patrick L., Corvallis, OR, United States AVI BioPharma, Inc., Corvallis, OR, United States (U.S. INVENTOR(S): PATENT ASSIGNEE(S):

corporation)

NUMBER KIND DATE us 6365577 PATENT INFORMATION: в1 20020402 APPLICATION INFO.: us 1999-426804 19991022 (9)

> NUMBER DATE

PRIORITY INFORMATION: US 1998-105695P 19981026 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED Wang, Andrew Zara, Jane PRIMARY EXAMINER: ASSISTANT EXAMINER: LEGAL REPRESENTATIVE: Gorthey, LeeAnn

NUMBER OF CLAIMS: 18

EXEMPLARY CLAIM: NUMBER OF DRAWINGS: 2 Drawing Figure(s); 2 Drawing Page(s)

LINE COUNT: 1006

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Antisense oligonucleotides useful for treating a disease state AB ***p53*** induction, such as proliferative cell characterized by disorders, e.g. cancer, or a hypoxic state induced by an ischemic attack, such as stroke, are described. The ***antisense*** are preferably of the class known as "steric blocker" type oligonucleotides, including ***morpholino*** oligonucleotides, peptide nucleic acids, 2'-O-allyl or 2'-O-alkyl modified oligonucleotides, or N3'.fwdarw.P5' phosphoramidate oligonucleotides.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 3 OF 41 CAPLUS COPYRIGHT 2003 ACS 2002:946579 CAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER: 138:21196

TITLE: Peroxisomal enoyl-CoA isomerases as modifiers of the

p53 pathway and their use in diagnosis and treatment

of p53-related diseases

INVENTOR(S): Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia;

Francis-Lang, Helen Exelixis, Inc., USA PCT Int. Appl., 44 pp. PATENT ASSIGNEE(S): SOURCE:

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English 37

FAMILY ACC. NUM. COUNT: PATENT INFORMATION:

> PATENT NO. KIND DATE APPLICATION NO. DATE wo 2002099426 20021212 wo 2002-us17420 20020603 Α1 AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,

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                           BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG
95 A1 20021219 US 2002-161510 20020603
                                                                       us 2002-161510
us 2002-161398
         us 2002192695
         us 2003013144
                                      Α1
                                               20030116
                                                                                                    20020603
                                                                   US 2001-296076P P
PRIORITY APPLN. INFO.:
                                                                                                    20010605
                                                                   US 2001-328605P P
                                                                                                  20011010
                                                                   US 2002-357253P P 20020215
        Two human peroxisomal .DELTA.3,.DELTA.2-enoyl-CoA isomerase (PECI) genes are identified as modulators of the p53 pathway, and thus are therapeutic targets for disorders assocd. with defective p53 function. Genetic
AB
         screens were designed to identify modifiers of the p53 pathway in
        Drosophila in which p53 was overexpressed in the wing. The CG13890 gene was identified as a modifier of the p53 pathway. Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, PECI
        genes and proteins are attractive drug targets for the treatment of
        pathologies assocd. with a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening for agents that modulate the activity of PECI are provided.

ZENCE COUNT: 2 THERE ARE 2 CITED REFERENCES AVAILABLE FOR THIS
REFERENCE COUNT:
                                                   RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT
L11 ANSWER 4 OF 41 CAPLUS COPYRIGHT 2003 ACS
                                          2002:946511 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                                          138:21195
TITLE:
                                         Glycine receptor chloride channel proteins as
                                         modifiers of the p53 pathway and their use in diagnosis and treatment of p53-related diseases Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia;
INVENTOR(S):
                                          Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
PATENT ASSIGNEE(S):
                                          Exelixis, Inc., USA
SOURCE:
                                         PCT Int. Appl., 56 pp.
                                          CODEN: PIXXD2
DOCUMENT TYPE:
                                         Patent
LANGUAGE:
                                         English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
        PATENT NO.
                                    KIND DATE
                                                                       APPLICATION NO.
                                                                                                   DATE
        wo 2002099140
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                                              20021212
                                                                       wo 2002-us17458 20020602
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                     GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR,
                     LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
                     TJ, TM
              RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH,
                     CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG US 2002-161510 20020603
        US 2002192695
                                                                       US 2002-161398
        US 2003013144
                                     Α1
                                              20030116
                                                                                                   20020603
                                                                  US 2001-296076P P
US 2001-328605P P
US 2002-357253P P
PRIORITY APPLN. INFO.:
                                                                                                   20010605
                                                                                                   20011010
                                                                                                  20020215
AΒ
        Three human glycine receptor chloride channel .alpha.-subunit (GLRA) genes
       are identified as modulators of the p53 pathway, and thus are therapeutic targets for disorders assocd. with defective p53 function. Genetic screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The CG14723 gene was identified as a modifier of the p53 pathway. Accordingly, vertex
        orthologs of these modifiers, and preferably the human orthologs, GLRA
       genes and proteins are attractive drug targets for the treatment of pathologies assocd. with a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening
        for agents that modulate the activity of GLRA are provided.
REFERENCE COUNT:
                                                   THERE ARE 4 CITED REFERENCES AVAILABLE FOR THIS
                                                   RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT
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L11 ANSWER 5 OF 41 CAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:946509 CAPLUS

DOCUMENT NUMBER:

138:21194

TITLE:

Human sulfotransferase proteins as modifiers of the p53 pathway and their use in diagnosis and treatment of p53-related diseases

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Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
                                    Exelixis, Inc., USA
PATENT ASSIGNEE(S):
SOURCE:
                                    PCT Int. Appl., 54 pp.
                                    CODEN: PIXXD2
DOCUMENT TYPE:
                                    Patent
                                    English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
       PATENT NO.
                               KIND DATE
                                                              APPLICATION NO.
                                                                                      DATE
       wo 2002099138
                                        20021212
                                                              wo 2002-us17409 20020603
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                   LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH,
                   PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
                   TJ, TM
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       us 2002192695
       us 2003013144
                                        20030116
                                                              us 2002-161398
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                                                          US 2001-296076P P
PRIORITY APPLN. INFO.:
                                                                                      20010605
                                                         US 2001-328605P P
US 2002-357253P P
                                                                                      20011010
                                                                                      20020215
       Four human sulfotransferase proteins with transferase domains (HS2ST)
AΒ
       genes are identified as modulators of the p53 pathway, and thus are
       therapeutic targets for disorders assocd. with defective p53 function.
       Genetic screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The pipe gene was identified as a modifier of the p53 pathway. Accordingly, vertebrate
       orthologs of these modifiers, and preferably the human orthologs, HS2ST
       genes and proteins are attractive drug targets for the treatment of pathologies assocd. with a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening
       for agents that modulate the activity of HS2ST are provided.
L11 ANSWER 6 OF 41 CAPLUS COPYRIGHT 2003 ACS
                                    2002:946461 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                                    138:35033
TITLE:
                                    cDNA and protein sequences of human glutamine
                                    fructose-6-phosphate amidotransferase and the uses of
the protein as modifiers of the p53 pathway in
                                    diagnosis and therapeutics
                                    Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia; Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
INVENTOR(S):
                                    Exelixis, Inc., USA PCT Int. Appl., 55 pp.
PATENT ASSIGNEE(S):
SOURCE:
                                    CODEN: PIXXD2
DOCUMENT TYPE:
                                    Patent
LANGUAGE:
                                    English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
       PATENT NO.
                               KIND DATE
                                                              APPLICATION NO.
                                                                                      DATE
       wo 2002099083
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                                        20021212
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                  PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
                   TJ, TM
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                       BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, 95 A1 20021219 US 2002-161510 20020603 44 A1 20030116 US 2002-161398 20020603
                  BF,
       US 2002192695
       US 2003013144
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US 2001-328605P P US 2002-357253P P 20020215 Human glutamine fructose-6-phosphate amidotransferase (GFAT) genes are AB identified as modulators of the p53 pathway, and thus are therapeutic

US 2001-296076P P

20010605

20011010

PRIORITY APPLN. INFO.:

screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The human GFAT gene was identified by Blast searching of mouse GFAT counterparts and the invention also provides distribution pattern of human GFAT gene in normal and tumor tissue cells. Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, GFAT genes and proteins are attractive drug targets for the treatment of pathologies assocd. With a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening for agents that modulate the activity of GFAT are provided.

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L11 ANSWER 7 OF 41 CAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:946453 CAPLUS
                                                138:35032
DOCUMENT NUMBER:
 TITLE:
                                                Protein arginine N-methyltransferase as modifiers of
                                                the p53 pathway and their use in diagnosis and
                                                treatment of p53-related diseases
                                                Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia; Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
 INVENTOR(S):
 PATENT ASSIGNEE(S):
                                                Exelixis, Inc., USA
 SOURCE:
                                                PCT Int. Appl., 67 pp.
                                                CODEN: PIXXD2
DOCUMENT TYPE:
                                                Patent
                                                English
LANGUAGE:
 FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
          PATENT NO.
                                         KIND DATE
                                                                                APPLICATION NO. DATE
         wo 2002099075
                                          A2 20021212
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         Human and mouse protein arginine N-methyltransferase (PRMT) genes are
         identified as modulators of the p53 pathway, and thus are therapeutic targets for disorders assocd. with defective p53 function. Genetic screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The C65358 gene was identified as a modifier of the p53 pathway. Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, PRMT genes and proteins are attractive drug targets for the treatment of pathologies assocd. With a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening for agents that modulate the activity of PRMT are provided.
         for agents that modulate the activity of PRMT are provided.
L11 ANSWER 8 OF 41 CAPLUS COPYRIGHT 2003 ACS
                                               2002:946452 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                                               138:34194
TITLE:
                                               cDNA and protein sequences of human amino acid
                                               transporter SLC7s and the uses of the protein as modifiers of the p53 pathway in diagnosis and
INVENTOR(S):
                                               Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia;
                                               Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
PATENT ASSIGNEE(S):
                                               Exelixis, Inc., USA PCT Int. Appl., 129 pp.
SOURCE:
                                               CODEN: PIXXD2
DOCUMENT TYPE:
                                               Patent
LANGUAGE:
                                               English
FAMILY ACC. NUM. COUNT: 37
PATENT INFORMATION:
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APPLICATION NO. DATE

PATENT NO.

KIND DATE

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US 2001-296076P P 20010015
PRIORITY APPLN. INFO.:
                                                               US 2001-328605P
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                                                                                              20011010
                                                               US 2001-338733P P
                                                                                              20011022
                                                               US 2002-357253P P
                                                                                              20020215
                                                               US 2002-357600P P
                                                                                              20020215
        Human amino acid transporter SLC7 genes are identified as modulators of
ΑB
        the p53 pathway, and thus are therapeutic targets for disorders assocd.
        with defective p53 function. Genetic screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed
        in the wing. The human SLC7 genes were identified by Blast searching of
       mouse SLC7 counterparts and the invention also provides distribution pattern of human SLC7 genes in normal and tumor tissue cells. Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, SLC7 genes and proteins are attractive drug targets for the treatment of pathologies assocd with a defective p53 signaling
        pathway, such as cancer. Methods for identifying modulators of p53
        comprising screening for agents that modulate the activity of SLC7 are
        provided.
L11 ANSWER 9 OF 41 CAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:946437 CAPLUS
                                       138:20565
DOCUMENT NUMBER:
TITLE:
                                       cDNA and protein sequences of human U5-snRNP-specific
                                       protein U5-200KD and the uses of the protein as
                                       modifiers of the p53 pathway in diagnosis and
                                       therapeutics
INVENTOR(S):
                                       Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia;
                                       Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
PATENT ASSIGNEE(S):
                                       Exelixis, Inc., USA
SOURCE:
                                       PCT Int. Appl., 87 pp.
                                       CODEN: PIXXD2
                                       Patent
DOCUMENT TYPE:
LANGUAGE:
                                       English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
        PATENT NO.
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                                           DATE
                                                                   APPLICATION NO.
                                                                                              DATE
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                                            20021212
                                                                   wo 2002-us17524
                                                                                              20020603
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                    GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR,
                    LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH,
                    PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
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        us 2002192695
        US 2003013144
                                            20030116
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                                    Α1
                                                                                              20020603
PRIORITY APPLN. INFO.:
                                                              US 2001-296076P P
US 2001-328605P P
                                                                                              20010605
                                                                                              20011010
                                                               US 2002-357253P
                                                                                              20020215
ΑB
       Human U5-snRNP-specific protein U5-200KD genes are identified as
       modulators of the p53 pathway, and thus are therapeutic targets for disorders assocd. with defective p53 function. Genetic screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The human U5-200KD genes were
       identified by Blast searching of mouse U5-200KD counterparts and the invention also provides distribution pattern of human U5-200KD genes in
       normal and tumor tissue cells. Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, U5-200KD genes and proteins
       are attractive drug targets for the treatment of pathologies assocd. with
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a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening for agents that

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L11 ANSWER 10 OF 41 CAPLUS COPYRIGHT 2003 ACS
                                2002:946436 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                                138:34193
                                cDNA and protein sequences of human potassium channel
TITLE:
                                and the uses of the protein as modifiers of the p53
                                pathway in diagnosis and therapeutics
                                Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia; Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
INVENTOR(S):
PATENT ASSIGNEE(S):
                                Exelixis, Inc., USA
                                PCT Int. Appl., 62 pp.
SOURCE:
                                CODEN: PIXXD2
DOCUMENT TYPE:
                                Patent
LANGUAGE:
                                English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
      PATENT NO.
                            KIND DATE
                                                      APPLICATION NO.
                                                                            DATE
      wo 2002099058
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                                   20021212
                                                      wo 2002-us17476 20020603
           W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
                CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH,
                PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
           BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG 92695 A1 20021219 US 2002-161510 20020603
      us 2002192695
                                                      US 2002-161398
      us 2003013144
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                                   20030116
                                                                            20020603
                                                   US 2001-296076P P
PRIORITY APPLN. INFO.:
                                                                            20010605
                                                   US 2001-328605P P
                                                                           20011010
                                                   US 2002-357253P P 20020215
AB
      Human potassium large conductance calcium-activated channel (subfamily M)
      alpha member (KCNMA) genes are identified as modulators of the p53 pathway, and thus are therapeutic targets for disorders assocd. With
      defective p53 function. Genetic screens were designed to identify
      modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The human KCNMA genes were identified by Blast searching of mouse KCNMA counterparts and the invention also provides distribution
      pattern of human KCNMA genes in normal and tumor tissue cells.
      Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, KCNMA genes and proteins are attractive drug targets for the treatment of pathologies assocd. With a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53,
      comprising screening for agents that modulate the activity of KCNMA are
      provided.
L11 ANSWER 11 OF 41 CAPLUS COPYRIGHT 2003 ACS
                               2002:946435 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                               138:21191
                               RNA-binding KH domain-containing proteins as modifiers of the p53 pathway and their use in diagnosis and
TITLE:
                               treatment of p53-related diseases
                               Friedman, Lori; Plowman, Gregory D.; Belvin, Marcia;
INVENTOR(S):
                               Francis-Lang, Helen; Li, Danxi; Funke, Roel P.
PATENT ASSIGNEE(S):
                               Exelixis, Inc., USA
                               PCT Int. Appl., 53 pp.
SOURCE:
                               CODEN: PIXXD2
DOCUMENT TYPE:
                               Patent
LANGUAGE:
                               English
FAMILY ACC. NUM. COUNT:
                               37
PATENT INFORMATION:
      PATENT NO.
                           KIND DATE
                                                      APPLICATION NO.
                                                                            DATE
      wo 2002099057
                            Α2
                                   20021212
                                                      WO 2002-US17475
                                                                            20020603
               AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
           W:
                CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
                UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
                TJ, TM
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CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG. 92695 A1 20021219 US 2002-161510 20020603
       us 2002192695
                                                                  us 2002-161510
       US 2003013144
                                                                  us 2002-161398
                                   Α1
                                           20030116
                                                                                             20020603
                                                              US 2001-296076P P
US 2001-328605P P
PRIORITY APPLN. INFO.:
                                                                                             20010605
                                                                                             20011010
                                                              US 2002-357253P P 20020215
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Four human RNA-binding proteins with KH domains (SAM) genes are identified as modulators of the p53 pathway, and thus are therapeutic targets for disorders assocd. with defective p53 function. Genetic screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The qkr58E-2 gene was identified as a modifier of the p53 pathway. Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, SAM genes and proteins are attractive drug targets for the treatment of pathologies assocd with a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening for agents that modulate the activity of SAM are provided.

L11 ANSWER 12 OF 41 CAPLUS COPYRIGHT 2003 ACS 2002:946434 CAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER: 138:20564

CDNA and protein sequences of human protein C22C7ORF and the uses of the protein as modifiers of the p53 TITLE:

pathway in diagnosis and therapeutics

INVENTOR(S): Friedman, Lori; Plowman, Gregory_D.; Belvin, Marcia;

Francis-Lang, Helen; Li, Danxi; Funke, Roel P.

Exelixis, Inc., USA PCT Int. Appl., 52 pp. PATENT ASSIGNEE(S): SOURCE:

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

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PATENT NO.
                             KIND DATE
                                                          APPLICATION NO.
                                                                                 DATE
      wo 2002099056
                              Α2
                                      20021212
                                                          wo 2002-us17474 20020603
            W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
                 CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
                 GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ,
                 UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
                 TJ., TM
            RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR,
                 BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG
92695 A1 20021219 US 2002-161510 20020603
      us 2002192695
      us 2003013144
                               Α1
                                      20030116
                                                          US 2002-161398
                                                                                  20020603
                                                      US 2001-296076P P
PRIORITY APPLN. INFO.:
                                                                                 20010605
                                                      US 2001-328605P P
                                                                                 20011010
                                                      US 2002-357253P P
                                                                                 20020215
```

AB Human protein C22C7ORF genes are identified as modulators of the p53 pathway, and thus are therapeutic targets for disorders assocd. with defective p53 function. Genetic screens were designed to identify modifiers of the p53 pathway in Drosophila in which p53 was overexpressed in the wing. The human protein C22C7ORF genes were identified by Blast searching of mouse protein C22C7ORF counterparts and the invention also provides distribution pattern of human protein C22C7ORF genes in normal and cancerous tissue cells. Accordingly, vertebrate orthologs of these modifiers, and preferably the human orthologs, protein C22C7ORF genes and proteins are attractive drug targets for the treatment of pathologies assocd. With a defective p53 signaling pathway, such as cancer. Methods for identifying modulators of p53, comprising screening for agents that modulate the activity of protein C22C7ORF are provided.

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L11 ANSWER 13 OF 41 USPATFULL
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ACCESSION NUMBER: 2002:337337 USPATFULL

TITLE:

PIBs as modifiers of the p53 pathway and methods of use Friedman, Lori, San Francisco, CA, UNITED STATES
Plowman, Gregory D., San Carlos, CA, UNITED STATES INVENTOR(S):

Belvin, Marcia, Albany, CA, UNITED STATES

Francis-Lang, Helen, San Francisco, CA, UNITED STATES

Li, Danxi, San Francisco, CA, UNITED STATES

Funke, Roel P., South San Francisco, CA, UNITED STATES

NUMBER KIND DATE PATENT INFORMATION: us 2002192695 20021219 Α1 US 2002-161510 APPLICATION INFO.: Α1 20020603 (10)

NUMBER DATE

PRIORITY INFORMATION:

US 2001-296076P 20010605 (60) US 2001-328605P 20011010 (60) 20020215 (60)

US 2002-357253P

DOCUMENT TYPE:

Utility APPLICATION

FILE SEGMENT: LEGAL REPRESENTATIVE:

JAN P. BRUNELLE, EXELIXIS, INC., 170 HARBOR WAY, P.O.

BOX 511, SOUTH SAN FRANCISCO, CA, 94083-0511

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

25 1

LINE COUNT:

4841

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Human PIB genes are identified as modulators of the p53 pathway, and thus are therapeutic targets for disorders associated with defective p53 function. Methods for identifying modulators of p53, comprising

screening for agents that modulate the activity of PIB are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 14 OF 41 USPATFULL

ACCESSION NUMBER:

2002:301576 USPATFULL

TITLE: INVENTOR(S): Inhibition of ATF2 activity to treat cancer Ronai, Ze?apos,ev, Suffern, NY, UNITED STATES Mount Sinai School of Medicine (U.S. corporation)

PATENT ASSIGNEE(S):

NUMBER **KIND** DATE PATENT INFORMATION: US 2002169121 20021114 Α1 us 2002-76905 APPLICATION INFO.: 20020214 (10)Α1

> NUMBER DATE

PRIORITY INFORMATION:

US 2001-269257P 20010216 (60) 20010215 (60)

US 2001-269118P Utility

DOCUMENT TYPE: FILE SEGMENT:

APPLICATION

LEGAL REPRESENTATIVE:

DARBY & DARBY P.C., 805 Third Avenue, New York, NY,

10022

NUMBER OF CLAIMS:

32

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS:

11 Drawing Page(s) 3146

LINE COUNT:

The present invention relates to novel therapies for cancer and, in particular, to therapies that are particularly suited to tumor cells resistant to other types of therapies such as radiation, chemotherapy, or combinations of both approaches. The invention provides methods for identifying and implementing strategies to inhibit a transcription factor which, in combination with other factors, renders the cells resistant and inhibits apopotosis of the cells. The invention provides an inhibitory ATF2 N-terminal fragment, specifically a fragment corresponding to amino acid residues 50-100 of ATF2 (termed peptide II). The invention provides methods for inhibiting tumor cell growth with such peptides.

L11 ANSWER 15 OF 41 USPATFULL

ACCESSION NUMBER:

2002:295143 USPATFULL

TITLE:

INVENTOR(S):

Oligoribonucleotides and ribonucleases for cleaving RNA

Crooke, Stanley T., Carlsbad, CA, UNITED STATES

NUMBER **KIND** DATE US 2002165189 20021107 Α1

PATENT INFORMATION: APPLICATION INFO.:

US 2002-78949 20020220 Α1 (10)

RELATED APPLN. INFO.:

Continuation of Ser. No. US 2000-479783, filed on 7 Jan

2000, PENDING

DOCUMENT TYPE: FILE SEGMENT:

Utility APPLICATION

LEGAL REPRESENTATIVE:

Woodcock Washburn LLP, One Liberty Place, 46th Floor,

Philadelphia, PA, 19103

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS:

10 Drawing Page(s)

LINE COUNT:

3922

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Oligomeric compounds including oligoribonucleotides and oligoribonucleosides are provided that have subsequences of 2'-pentoribofuranosyl nucleosides that activate dsRNase. The oligoribonucleotides and oligoribonucleosides can include substituent groups for increasing binding affinity to complementary nucleic acid strand as well as substituent groups for increasing nuclease resistance. The oligomeric compounds are useful for diagnostics and other research purposes, for modulating the expression of a protein in organisms, and for the diagnosis, detection and treatment of other conditions susceptible to oligonucleotide therapeutics. Also included in the invention are mammalian ribonucleases, i.e., enzymes that degrade RNA, and substrates for such ribonucleases. Such a ribonuclease is referred and substrates for such ribonucleases. Such a ribonuclease is referred to herein as a dsRNase, wherein "ds" indicates the RNase's specificity for certain double-stranded RNA substrates. The artificial substrates for the dsRNases described herein are useful in preparing affinity matrices for purifying mammalian ribonuclease as well as non-degradative RNA-binding proteins.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 16 OF 41 USPATFULL

ACCESSION NUMBER:

TITLE:

2002:294546 USPATFULL
Detection of nucleic acid heteroduplex molecules by

anion-exchange chromatography

INVENTOR(S):

Taylor, Paul D., Gilroy, CA, UNITED STATES

NUMBER KIND DATE US 2002164589 20021107 A1

PATENT INFORMATION: APPLICATION INFO.:

Α1 US 2001-756070 20010106 (9)

RELATED APPLN. INFO.:

Continuation-in-part of Ser. No. US 2000-687834, filed

on 11 Oct 2000, PENDING

NUMBER DATE

PRIORITY INFORMATION:

Utility

US 2000-194652P 20000404 (60)

DOCUMENT TYPE: FILE SEGMENT:

APPLICATION

LEGAL REPRESENTATIVE:

JOHN F. BRADY, TRANSGENOMIC, INC., 2032 CONCOURSE

DRIVE, SAN JOSE, CA, 95131

NUMBER OF CLAIMS:

EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

1 Drawing Page(s)

LINE COUNT: 2151

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

72

The present invention describes a method for separating or partially separating heteroduplex and homoduplex DNA molecules in a mixture. In the method, the mixture is applied to an anion-exchange chromatography medium. The heteroduplex and homoduplex molecules are eluted with a medium. The neteroduplex and nomoduplex molecules are eluted with a mobile phase containing an eluting salt, including an anion and a cation, a buffer, and preferably including an organic solvent. The eluting is carried out under conditions effective to at least partially denature the heteroduplexes (e.g., thermal or chemical denaturing) resulting in the separation of the heteroduplexes from the homoduplexes. The method has many applications including, but not limited to, detecting mutations and comparative DNA sequencing.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 17 OF 41 USPATFULL

ACCESSION NUMBER:

2002:273383 USPATFULL

TITLE:

Antisense oligonucleotide modulation of human MDM2

INVENTOR(S):

expression Miraglia, Loren J., Encinitas, CA, UNITED STATES

Nero, Pamela, Oceanside, CA, UNITED STATES Graham, Mark J., San Clemente, CA, UNITED STATES Monia, Brett P., La Costa, CA, UNITED STATES

NUMBER KIND DATE US 2002151511 20021017 Α1 US 2001-851771 Α1 20010509 (9)

PATENT INFORMATION: APPLICATION INFO.:

1998, GRANTED, Pat. No. US 6238921

Utility DOCUMENT TYPE: FILE SEGMENT: APPLICATION

Licata & Tyrrell P.C., 66 E. Main Street, Marlton, NJ, LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 40 **EXEMPLARY CLAIM:** 1 1409 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Compounds, compositions and methods are provided for inhibiting the expression of human mdm2. The compositions comprise antisense

oligonucleotides targeted to nucleic acids encoding mdm2. Methods of using these oligonucleotides for inhibition of mdm2 expression and for treatment of diseases such as cancers associated with overexpression of

mdm2 are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 18 OF 41 USPATFULL

2002:272813 USPATFULL ACCESSION NUMBER:

Diagnosis and treatment of cancer using mammalian TITLE:

pellino polypeptides and polynucleotides

INVENTOR(S): Powers, Scott, Greenlawn, NY, UNITED STATES

Mu, David, Jericho, NY, UNITED STATES Xiang, Phil, San Francisco, CA, UNITED STATES

Peng, Yue, South Setauket, NY, UNITED STATES

Tularik Inc., South San Francisco, CA, 94080 (U.S. PATENT ASSIGNEE(S):

corporation)

NUMBER KIND DATE Α1 20021017

US 2002150934 US 2001-41030 PATENT INFORMATION: APPLICATION INFO.: Α1 20011228 (10)

> NUMBER DATE

US 2001-259502P 20010102 (60)

PRIORITY INFORMATION: DOCUMENT TYPE: Utility

FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: TOWNSEND AND TOWNSEND AND CREW, LLP, TWO EMBARCADERO

CENTER, EIGHTH FLOOR, SAN FRANCISCO, CA, 94111-3834

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 3 Drawing Page(s)

LINE COUNT: 3106

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides methods, reagents, and kits for diagnosing and treating cancer in a mammal, e.g., a human. This invention is based upon the discovery that Pellino 1 or 2 is overexpressed and/or amplified in cancer. Methods to detect cancer or a

propensity to develop cancer, to monitor the efficacy of a cancer treatment, and to treat cancer, by inhibiting the expression and/or activity of Pellino 1 or 2 in a cancer cell are included.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 19 OF 41 USPATFULL

ACCESSION NUMBER: 2002:266324 USPATFULL Tyrosine kinase inhibitors TITLE:

INVENTOR(S): Bilodeau, Mark T., Lansdale, PA, UNITED STATES Hartman, George D., Lansdale, PA, UNITED STATES Manley, Peter J., Harleysville, PA, UNITED STATES

PATENT ASSIGNEE(S): Merck & Co., Inc. (U.S. corporation)

NUMBER KIND DATE

US 2002147203 PATENT INFORMATION: Δ1 20021010 APPLICATION INFO.: us 2002-62351 Α1 20020201 (10)

RELATED APPLN. INFO.: Continuation of Ser. No. US 2000-658680, filed on 8 Sep

2000, ABANDONED

NUMBER DATE

PRIORITY INFORMATION:

US 1999-153348P 19990910 (60)

DOCUMENT TYPE: FILE SEGMENT:

Utility **APPLICATION** RY60-30, Rahway, NJ, 07065-0907

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1 3989 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to compounds which inhibit, regulate and/or modulate tyrosine kinase signal transduction, compositions which contain these compounds, and methods of using them to treat tyrosine kinase-dependent diseases and conditions, such as angiogenesis, cancer, tumor growth, atherosclerosis, age related macular degeneration, diabetic retinopathy, inflammatory diseases, and the like in mammals.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 20 OF 41 USPATFULL

ACCESSION NUMBER: 2002:266257 USPATFULL

TITLE:

Compounds for targeting endothelial cells, compositions containing the same and methods for their use Von Wronski, Mathew A., Moorestown, NJ, UNITED STATES INVENTOR(S):

Marinelli, Edmund R., Lawrenceville, NJ, UNITED STATES Nunn, Adrian D., Lambertville, NJ, UNITED STATES Pillai, Radhakrishna, Cranbury, NJ, UNITED STATES Ramalingam, Kondareddiar, Dayton, NJ, UNITED STATES Tweedle, Michael F., Princeton, NJ, UNITED STATES

Linder, Karen, Kingston, NJ, UNITED STATES

Nanjappan, Palaniappa, Dayton, NJ, UNITED STATES Raju, Natarajan, Kendall Park, NJ, UNITED STATES

NUMBER KTND DATE US 2002147136 20021010 Α1

PATENT INFORMATION: APPLICATION INFO.: US 2001-871974 Α1 20010604 (9)

Continuation-in-part of Ser. No. US 2000-585364, filed RELATED APPLN. INFO.:

on 2 Jun 2000, PENDING

DOCUMENT TYPE: Utility FILE SEGMENT: **APPLICATION**

NIXON & VANDERHYE P.C., 8th Floor, 1100 North Glebe Road, Arlington, VA, 22201-4714 LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 4 Drawing Page(s)

LINE COUNT: 5017

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides compounds for targeting endothelial cells, tumor cells or other cells that express the NP-1 receptor,

compositions containing the same and methods for their use. Additionally, the present invention includes diagnostic, therapeutic and radiotherapeutic compositions useful for visualization, therapy or

radiotherapy.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 21 OF 41 USPATFULL

ACCESSION NUMBER: 2002:165211 USPATFULL

Use of Rad51 inhibitors for p53 gene therapy TITLE: Zarling, David A., Menlo Park, CA, UNITED STATES Reddy, Gurucharan, Fremont, CA, UNITED STATES INVENTOR(S):

NUMBER KIND DATE PATENT INFORMATION: US 2002086840 20020704 Α1 APPLICATION INFO.: US 2001-771355 Α1 20010126 (9)

> DATE NUMBER

PRIORITY INFORMATION: US 2000-178561P 20000126 (60)

DOCUMENT TYPE: Utility

FILE SEGMENT: APPLICATION

FLEHR HOHBACH TEST, ALBRITTON & HERBERT LLP, Four Embarcarero Center, Suite 3400, San Francisco, CA, LEGAL REPRESENTATIVE:

94111-4187

NUMBER OF CLAIMS: 18 **EXEMPLARY CLAIM:** LINE COUNT: 995

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention is directed to methods and compositions for

vivo. More specifically, a tumor cell is contacted, in vivo, with a Rad51 inhibitor, and a polynucleotide capable of expressing functional ***p53*** protein. In a further embodiment of the present invention the tumor cell is exposed in vivo to radiation or chemotherapeutic agents (e.g., BCNU, CCNU, and DMZ, GB, cisplatin and the like). The Rad51 inhibitor may be selected from the group consisting of peptides, small molecules and Rad51 ***antisense*** molecules. The Rad51 ***antisense*** polynucleotide may be accorded on an expression vector under the control of one or more encoded on an expression vector under the control of one or more promoters, and the expression vector may then be incorporated into a viral genome, preferably an andeno or retro virus, which is then used to introduce the expression vector into the tumor cell.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 22 OF 41 USPATFULL

ACCESSION NUMBER: 2002:141539 USPATFULL

TITLE: INVENTOR(S):

Orally active salts with tyrosine kinase activity Fraley, Mark E., North Wales, PA, UNITED STATES Karki, Shyam B., Lansdale, PA, UNITED STATES Kim, Yuntae, Harleysville, PA, UNITED STATES

NUMBER KIND DATE

PATENT INFORMATION: APPLICATION INFO.:

US 2002072526 20020613 Α1 us 2001-981979 (9) Α1 20011017

> DATE NUMBER

PRIORITY INFORMATION:

US 2000-241043P 20001017 (60)

DOCUMENT TYPE:

Utility APPLICATION

FILE SEGMENT: LEGAL REPRESENTATIVE:

MERCK AND CO INC, P O BOX 2000, RAHWAY, NJ, 070650907

NUMBER OF CLAIMS:

39

EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

3 Drawing Page(s)

LINE COUNT:

1971 CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB

The present invention relates to orally active salts of compounds which inhibit, regulate and/or modulate tyrosine kinase signal transduction, compositions which contain these compounds, and methods of using them to treat tyrosine kinase-dependent diseases and conditions, such as angio-genesis, cancer, tumor growth, atherosclerosis, age related macular degeneration, diabetic retinopathy, inflammatory diseases, and the like in mammals.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 23 OF 41 USPATFULL

ACCESSION NUMBER:

2002:297601 USPATFULL

TITLE:

Tyrosine kinase inhibitors

INVENTOR(S):

Fraley, Mark E., North Wales, PA, United States Hambaugh, Scott R., Norristown, PA, United States Hungate, Randall W., Lansdale, PA, United States

PATENT ASSIGNEE(S):

Merck & Co., Inc., Rahway, NJ, United States (U.S.

corporation)

NUMBER KIND DATE 20021112 в1

PATENT INFORMATION: APPLICATION INFO.:

US 6479512 US 2000-690602 20001017 (9)

> NUMBER DATE

PRIORITY INFORMATION: DOCUMENT TYPE:

19991019 (60)

US 1999-160362P Utility

FILE SEGMENT: GRANTED

PRIMARY EXAMINER:

LEGAL REPRESENTATIVE:

Seaman, D. Margaret Brown, Dianne, Daniel, Mark R.

NUMBER OF CLAIMS:

31

EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

0 Drawing Figure(s); 0 Drawing Page(s)

LINE COUNT:

2602

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB

The present invention relates to compounds which inhibit, regulate and/or modulate tyrosine kinase signal transduction, compositions which kinase-dependent diseases and conditions, such as angiogenesis, cancer, tumor growth, atherosclerosis, age related macular degeneration, diabetic retinopathy, inflammatory diseases, and the like in mammals.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 24 OF 41 USPATFULL

ACCESSION NUMBER:

2002:108831 USPATFULL TITLE:

ATM kinase modulation for screening and therapies Kastan, Michael, Cordova, TN, United States INVENTOR(S): Canman, Christine, Cordova, TN, United States Kim, Seong-Tae, Cordova, TN, United States Lim, Dae-Sik, Cordova, TN, United States St. Jude Children's Research Hospital, Memphis, TN, United States (U.S. corporation)

PATENT ASSIGNEE(S):

Johns-Hopkins University, Baltimore, MD, United States

(U.S. corporation)

NUMBER KIND DATE PATENT INFORMATION: US 6387640 20020514 в1

APPLICATION INFO.: DOCUMENT TYPE:

US 1999-248061 19990210 (9) Utility

FILE SEGMENT: GRANTED PRIMARY EXAMINER:

Achutamurthy, Ponnathapu

ASSISTANT EXAMINER: Monshipouri, M. LEGAL REPRESENTATIVE: Darby & Darby NUMBER OF CLAIMS:

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 0 Drawing Figure(s): 0 Drawing Page(s)

LINE COUNT: 2258

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to identification of the consensus sequence phosphorylated by ATM kinase. This, in turn, permitted identification of ATM kinase target proteins, and development of a convenient assay system for ATM kinase phosphorylation using fusion polypeptides as substrates. The assay system is adaptable to screening for ATM modulators, particularly inhibitors. In a specific embodiment, the substrate recognition sequence and mutagenized variants of this sequence were incorporated in a CET fusion protein and account for sequence were incorporated in a GST fusion protein and assayed for phosphorylation by ATM kinase. This assay system is useful in screening for ATM inhibitors. ATM function assays were validated using an ATM-kinase dead dominant-negative mutant.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 25 OF 41 USPATFULL

ACCESSION NUMBER: 2002:69773 USPATFULL

TITLE: Non-invasive method for detecting target RNA INVENTOR(S):

Iversen, Patrick L., Corvallis, OR, United States AVI BioPharma, Inc., Corvallis, OR, United States (U.S. PATENT ASSIGNEE(S):

corporation)

NUMBER KIND DATE PATENT INFORMATION: US 6365351 20020402 в1 us 2000-493494 APPLICATION INFO.: 20000128 (9)

NUMBER DATE

PRIORITY INFORMATION: US 1999-117846P 19990129 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED PRIMARY EXAMINER:

Wang, Andrew Zara, Jane ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Judge, Linda R., Perkins Coie LLP

NUMBER OF CLAIMS:

EXEMPLARY CLAIM: NUMBER OF DRAWINGS: 11 Drawing Figure(s); 3 Drawing Page(s) 1258 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ΑB The present invention provides a method for targeting a particular mRNA sequence in vivo by oral administration of a ***morpholino*** antisense compound having uncharged phosphorus-containing backbone linkages. Also disclosed is a non-invasive method of detecting and quantitating the in vivo presence of RNA containing one or more selected

nuclease-resistant antisense oligomer which hybridizes by Watson-Crick base pairing to a region of the target RNA with a Tm substantially greater than 37 degree. C. The oligomer is able to complex intracellularly with target RNA, and is released from intracellular sites as a nuclease-resistant heteroduplex, which can then be measured in a body fluid sample, e.g., urine.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 26 OF 41 USPATFULL

2002:34294 USPATFULL ACCESSION NUMBER:

ATM kinase modulation for screening and therapies TITLE:

INVENTOR(S):

Kastan, Michael, Cordova, TN, United States Canman, Christine, Cordova, TN, United States Kim, Seong-Tae, Cordova, TN, United States

PATENT ASSIGNEE(S):

Lim, Dae-Sik, Cordova, TN, United States
St. Jude Childre's Research Hospital, Memphis, TN,
United States (U.S. corporation)
Johns-Hopkins University, Baltimore, MD, United States

(U.S. corporation)

NUMBER KIND DATE PATENT INFORMATION:

US 6348311 20020219 в1 us 1999-400653 APPLICATION INFO.: 19990921

Continuation of Ser. No. US 1999-248061, filed on 10 RELATED APPLN. INFO.:

Feb 1999 Utility DOCUMENT TYPE: FILE SEGMENT: GRANTED

Prouty, Rebecca E. PRIMARY EXAMINER: Monshipouri, Maryam ASSISTANT EXAMINER: Darby & Darby

LEGAL REPRESENTATIVE: NUMBER OF CLAIMS:

EXEMPLARY CLAIM: NUMBER OF DRAWINGS: 9 Drawing Figure(s); 4 Drawing Page(s)

LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to identification of the consensus sequence phosphorylated by ATM kinase. This, in turn, permitted identification of ATM kinase target proteins, and development of a convenient assay system for ATM kinase phosphorylation using fusion polypeptides as substrates. The assay system is adaptable to screening for ATM modulators, particularly inhibitors. In a specific embodiment, the substrate recognition sequence and mutagenized variants of this sequence were incorporated in a GST fusion protein and assayed for phosphorylation by ATM kinase. This assay system is useful in screening for ATM inhibitors. ATM function assays were validated using an ATM-kinase dead dominant-negative mutant.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 27 OF 41 **MEDLINE DUPLICATE 2**

2002212307 ACCESSION NUMBER: MEDLINE

DOCUMENT NUMBER:

21945711 PubMed ID: 11948540 Bioavailability and efficacy of antisense TITLE:

morpholino oligomers targeted to c-myc and

cytochrome P-450 3A2 following oral administration in rats.

AUTHOR: Arora Vikram; Knapp Derek C; Ředdy Muralimohan T; Weller

Dwight D; Iversen Patrick L

AVI BioPharma, 4575 SW Research Way, Suite 200, Corvallis, CORPORATE SOURCE:

Oregon 97333, USA.. varora@avibio.com

GM54871 (NIGMS) CONTRACT NUMBER:

SOURCE: JOURNAL OF PHARMACEUTICAL SCIENCES, (2002 Apr) 91 (4)

1009-18.

Journal code: 2985195R. ISSN: 0022-3549.

PUB. COUNTRY:

United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE) LANGUAGE: English

Priority Journals FILE SEGMENT:

ENTRY MONTH: 200208

ENTRY DATE:

Entered STN: 20020412 Last Updated on STN: 20020815 Entered Medline: 20020814

Antisense ***Morpholino*** ΑB phosphorodiamidate oligomers (PMO) are resistant to degradation by cellular hydrolases, DNases, RNases, and phosphodiesterases, but remain sensitive to prolonged exposure to low

stability, and efficacy of two distinct PMO sequences targeted to c-myc and cytochrome P-450 (CYP) 3A2. The c-myc ***antisense*** 20-mer, AVI-4126 (5'-ACGTTGAGGGGCATCGTCGC-3'), slowed the regenerative process in the rat liver after a 70% partial hepatectomy (PH). Rats were administered 3.0 mg/kg AVI-4126 in 0.1 mL saline via a bolus intravenous injection or in 0.5 mL sterile phosphate-buffered saline via gavage immediately following PH. The areas under the plasma concentration versus time curves revealed a fractional oral availability of 78.8% over a period of 10 min through 24 h. Immunoblot analysis of liver tissue from rats treated orally with AVI-4126 demonstrated a sequence-specific reduction in the target protein c-Myc, as well as secondary proliferation markers: proliferating cell nuclear antigen (PCNA), cyclin D1, and ***p53*** . The CYP3A2

antisense 22-mer AVI-4472 (5'-GAGCTGAAAGCAGGTCCATCCC-3') caused a sequence-dependent reduction of approximately five-fold in the rat liver CYP3A2 protein levels and erythromycin demethylation activity in 24 h following oral administration at a dose of 2 mg/kg. It is concluded that oral administration of PMOs can inhibit c-myc and CYP3A2 gene expression in rat liver by an ***antisense*** -based mechanism of action. These studies highlight the potential for development of PMOs as orally administered therapeutic agents. Copyright 2002 Wiley-Liss, Inc. and the American Pharmaceutical Association J Pharm Sci 91:1009-1018, 2002

L11 ANSWER 28 OF 41 CAPLUS COPYRIGHT 2003 ACS

ACCESSION NUMBER:

2002:712518 CAPLUS

DOCUMENT NUMBER:

138:34532

TITLE: AUTHOR(S):

SOURCE:

Responses of human cells to PAH-induced DNA damage Baird, William M.; Hooven, Louisa A.; Mahadevan, Brinda; Luch, Andreas; Seidel, Albrecht; Iversen,

CORPORATE SOURCE:

Oregon State University, Corvallis, OR, 97331, USA Polycyclic Aromatic Compounds (2002), 22(3-4), 771-780 CODEN: PARCEO; ISSN: 1040-6638 Taylor & Francis Ltd.

PUBLISHER:

DOCUMENT TYPE:

Journal

LANGUAGE:

English Benzo[a]pyrene (B[a]P) and dibenzo[a,l]pyrene (DB[a,l]P) induce cytochrome P 450 (CYP) CYP1A1 and CYP1B1, which metabolize these polycyclic arom. hydrocarbons (PAHs) into DNA-binding species. In order to detail roles of CYPIA1 and CYPIB1 in activation of DB[a,1]P to the diol epoxide, we here report the inhibition of CYPIA1 in human MCF-7 cells with phosphorodiamidate ***morpholino*** ***antisense*** oligomers (morpholinos). PAH-DNA adduct formation was also detd. after treatment with morpholinos and B[a]P or DB[a,1]P. ***P53*** is involved in DNA is involved in DNA repair, cell cycle arrest, and apoptosis. ***p53*** Cells with normal protein arrest in the G1 phase of the cell cycle on exposure to DNA-damaging agents (presumably allowing the cell sufficient time to repair damaged DNA prior to replication). Previous studies in human Previous studies in human MCF-7 cells indicate that cells with PAH-DNA adducts escape cell cycle arrest and accumulate in the S phase. In the present study the effect of PAH-DNA adducts on the cell cycle were obsd. in human diploid fibroblasts (HDF). We found that treatment of HDF with the diol epoxide of DB[a,1]P causes cell cycle arrest in G1. An increase in DNA adduct formation with increase in concn. of dibenzo[a,l]pyrene diol epoxide {(-)-anti-DB[a,1]PDE} was also obsd.

REFERENCE COUNT: 36 THERE ARE 36 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L11 ANSWER 29 OF 41 **MEDLINE** DUPLICATE 3

ACCESSION NUMBER:

2002276603 MEDLINE

DOCUMENT NUMBER:

TITLE:

AUTHOR:

22011825 PubMed ID: 12015968 A dominant-negative form of p63 is required for epidermal

proliferation in zebrafish.

CORPORATE SOURCE:

Lee Hyunsook; Kimelman David
Department of Biochemistry, University of Washington, Box 357350, Seattle, WA 98195, USA.
Dev Cell, (2002 May) 2 (5) 607-16.
Journal code: 101120028. ISSN: 1534-5807. SOURCE:

PUB. COUNTRY:

DOCUMENT TYPE:

United States
Journal; Article; (JOURNAL ARTICLE)

English LANGUAGE:

FILE SEGMENT: Priority Journals OTHER SOURCE: GENBANK-AF487944 200206

ENTRY MONTH:

ENTRY DATE: Entered STN: 20020518

Last Updated on STN: 20020618

Epidermal stem cells play a critical role in producing the multilayered vertebrate skin. Products of the p63 gene not only mark the epidermal stem AB cells, but also are absolutely required for the formation of mammalian epidermis. We find that early zebrafish embryos express a dominant-negative form of p63 (DeltaNp63), which accumulates in the nucleus just as epidermal growth begins. Using ***antisense***

morpholino oligonucleotides, we show that DeltaNp63 is neg ***morpholino*** oligonucleotides, we show that DeltaNp63 is needed for epidermal growth and limb development and is specifically required for the proliferation of epidermal cells by inhibiting ***p53*** activity. While the structure of fish epidermis is very different from that of

higher vertebrates, our study shows that DeltaNp63 has essential and

L11 ANSWER 30 OF 41 MEDLINE **DUPLICATE 4**

ACCESSION NUMBER: 2002628967 MEDLINE

PubMed ID: 12386920 DOCUMENT NUMBER: 22274651

ancient role in the development of skin.

TITLE: Inhibition of human chorionic gonadotropin beta-subunit

modulates the mitogenic effect of c-myc in human prostate

cancer cells.

Devi Gayathri R; Oldenkamp Jennifer R; London Carla A; **AUTHOR:**

Iversen Patrick L

AVI BioPharma, Corvallis, Oregon 97333, USA.. grdevi@avibio.com CORPORATE SOURCE:

PROSTATE, (2002 Nov 1) 53 (3) 200-10. Journal code: 8101368. ISSN: 0270-4137. SOURCE:

PUB. COUNTRY: **United States**

Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

LANGUAGE: English

Priority Journals FILE SEGMENT:

ENTRY MONTH: 200211

ENTRY DATE: Entered STN: 20021019

Last Updated on STN: 20021213
Entered Medline: 20021121

BACKGROUND: Amplification of the proto-oncogene c-myc has been identified AB as one of the most common genetic alterations in prostate cancer, thus making it an attractive therapeutic target. However, certain prostate making it an attractive therapeutic target. However, certain product cancer cells are unresponsive to c-Myc inhibition. The purpose of this study was to test the hypothesis that effective growth inhibition in the refractory cancer cells can be achieved by blocking c-myc along with a growth factor using a novel phosphorodiamidate ***morpholino*** ***antisense*** oligomer-based approach. Human chorionic gonadotropin, a growth factor implicated in neoplasm, causes activation of c-myc through a G-protein-coupled pathway of signal transduction. METHODS: In this study, the effect of inhibition of beta-hCG and c-myc singly or in combination was evaluated in DU145 (RB -/-, ***p53*** -/-, androgen-independent) was evaluated in DU145 (RB -/-, and LNCaP (Rb+/+, ***p53*** and LNCaP (Rb+/+, ***p53*** +/+, androgen-sensitive) human prostate cancer cell lines and in a DU145 subcutaneous xenograft murine model.

RESULTS: ***Antisense*** phosphorodiamidate ***morpholino*** oligomers directed against beta-hCG and c-myc caused a specific decrease of the target protein levels. Unlike LNCaP cells, DU145 cell growth was refractory to c-Myc inhibition. Unresponsiveness to c-myc inhibition in DU145 cells was overcome by targeting both beta-hCG and c-myc genes, resulting in potentiation of the antiproliferative effect seen with inhibition of beta-hCG alone. CONCLUSIONS: The inhibition of beta-hCG sensitizes prostate cancer cells to the antiproliferative effects of c-Myc inhibition, including tumors that are refractory to c-Myc decrease alone. Copyright 2002 Wiley-Liss, Inc.

L11 ANSWER 31 OF 41 CAPLUS COPYRIGHT 2003 ACS

ACCESSION NUMBER: 2001:816897 CAPLUS

DOCUMENT NUMBER: 135:353717

TITLE: Splice-region antisense oligonucleotide composition

and targeting the mRNA splicing

INVENTOR(S): Iversen, Patrick L.; Hudziak, Robert PATENT ASSIGNEE(S):

Avi Biopharma, Inc., USA SOURCE: PCT Int. Appl., 53 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO. KIND DATE APPLICATION NO. DATE wo 2001083740 Α2 20011108 wo 2001-us14410 20010504 W: AU, CA, JP, KR

PT, SE, TR

PRIORITY APPLN. INFO.: US 2000-202376P P 20000504 Antisense compns. targeted against an mRNA sequence for a selected protein, at a region having its 5' end from 1 to about 25 base pairs downstream of a normal splice acceptor junction in the preprocessed mRNA, are disclosed. The antisense compd. is Rnase-inactive, and is preferably a phosphorodiamidate-linked ***morpholino*** oligonucleotide. Such targeting is effective to inhibit natural mRNA splice processing, produce splice variant mRNAs, and inhibit normal expression of the protein.

L11 ANSWER 32 OF 41 USPATFULL

2001:212454 USPATFULL ACCESSION NUMBER:

Tyrosine kinase inhibitors TITLE: INVENTOR(S):

Fraley, Mark E., North Wales, PA, United States Hartman, George D., Lansdale, PA, United States Hungate, Randall W., Newbury Park, CA, United States

NUMBER KIND DATE

PATENT INFORMATION: US 2001044451 Α1 20011122

us 6420382 В2 20020716 US 2001-788718 20010220 APPLICATION INFO.: Α1 (9)

NUMBER DATE

PRIORITY INFORMATION: US 2000-185023P 20000225 (60)

Utility DOCUMENT TYPE: FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: MERCK AND CO INC, P O BOX 2000, RAHWAY, NJ, 070650907

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1 2114 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to compounds which inhibit, regulate and/or modulate tyrosine kinase signal transduction, compositions which contain these compounds, and methods of using them to treat tyrosine kinase-dependent diseases and conditions, such as angiogenesis, cancer, tumor growth, atherosclerosis, age related macular degeneration, diabetic retinopathy, inflammatory diseases, and the like in mammals.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 33 OF 41 USPATFULL

ACCESSION NUMBER: 2001:165573 USPATFULL

TITLE: Non-invasive method for detecting target RNA INVENTOR(S):

Iversen, Patrick L., Corvallis, OR, United States AVI BioPharma, Inc. (U.S. corporation) PATENT ASSIGNEE(S):

NUMBER KIND DATE PATENT INFORMATION: US 2001024783 20010927 Α1 APPLICATION INFO.: US 2000-736920 Α1 20001213

Continuation-in-part of Ser. No. US 2000-493494, filed RELATED APPLN. INFO.:

(9)

on 28 Jan 2000, PENDING

NUMBER DATE

PRIORITY INFORMATION: US 1999-117846P 19990129 (60)

Utility DOCUMENT TYPE: FILE SEGMENT: **APPLICATION**

LEGAL REPRESENTATIVE: IOTA PI LAW GROUP, 350 CAMBRIDGE AVENUE SUITE 250, P O

BOX 60850, PALO ALTO, CA, 94306-0850

NUMBER OF CLAIMS: 31 EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

9 Drawing Page(s) LINE COUNT: 2004

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A method of detecting in a subject, the occurrence of a base-specific intracellular binding event involving a single-stranded target RNA, is disclosed. The method includes administering to the subject an oligomeric antisense compound having (i) from 8 to 40 bases, including a targeting base sequence that is complementary to a portion of the target RNA, (ii) a Tm, with respect to binding to a complementary RNA sequence, of greater than about 50.degree. C., and (iii) an ability to be actively taken up by mammalian cells, and (iv) conferring resistance of complementary RNA hybridized with the agent to RnaseH. Where the

substantially backbone. At a selected time after said administering the agent, a sample of a body fluid is obtained from the subject, and the presence in the sample of a nuclease-resistant heteroduplex composed of the antisense oligomer and the complementary portion of the target RNA is detected. The method is useful, for example, for detecting levels of gene expression, biochemical or physiological states that are characterized by expression of certain genes, genetic mutations, and the presence and identity of infective viral or bacterial agents. Also disclosed are arrays, kits and antibodies employed in carrying out the method.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 34 OF 41 USPATFULL

2001:139533 USPATFULL ACCESSION NUMBER:

Antisense modulation of human MDM2 expression TITLE: Miraglia, Loren J., Encinitas, CA, United States INVENTOR(S):

Nero, Pamela, Oceanside, CA, United States Graham, Mark J., San Clemente, CA, United States Monia, Brett P., La Costa, CA, United States Cowsert, Lex M., Carlsbad, CA, United States

KIND DATE NUMBER

us 2001016575 20010823 PATENT INFORMATION: Α1 APPLICATION INFO.: us 2001-752983 Α1 20010102

Continuation of Ser. No. US 1999-280805, filed on 26 RELATED APPLN. INFO.:

Mar 1999, GRANTED, Pat. No. US 6184212 Continuation-in-part of Ser. No. US 1998-48810, filed on 26 Mar 1998, GRANTED, Pat. No. US 6238921

Utility DOCUMENT TYPE: APPLICATION FILE SEGMENT:

LEGAL REPRESENTATIVE: Licata & Tyrrell P.C., 66 E. Main Street, Marlton, NJ,

08053

41 NUMBER OF CLAIMS: EXEMPLARY CLAIM: LINE COUNT: 3562

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Compounds, compositions and methods are provided for inhibiting the expression of human mdm2. The compositions include antisense compounds

targeted to nucleic acids encoding mdm2. Methods of using these

oligonucleotides for inhibition of mdm2 expression and for treatment of

diseases such as cancers associated with overexpression of mdm2 are

provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 35 OF 41 USPATFULL

ACCESSION NUMBER: 2001:215034 USPATFULL

TITLE: Arteriovenous and venous graft treatments: methods and

compositions

INVENTOR(S):

Zalewski, Andrew, Elkins Park, PA, United States Shi, Yi, Cheltenham, PA, United States Thomas Jefferson University, Philadelphia, PA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 6323184 20011127 в1 US 1995-424991 19950419 (8) APPLICATION INFO.:

RELATED APPLN. INFO.:

Continuation-in-part of Ser. No. wO 1994-US11853, filed on 17 Oct 1994 Continuation-in-part of Ser. No. US 1993-138637, filed on 15 Oct 1993, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: **GRANTED**

PRIMARY EXAMINER:

McGarry, Sean Drinker Biddle & Reath LLP LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 29 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 58 Drawing Figure(s); 31 Drawing Page(s)

LINE COUNT: 2692

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A method and compounds are provided for inhibiting the synthesis of extracellular matrix proteins. Compounds of the invention comprise oligonucleotides specific for nuclear proto-oncogenes. Preferably, oligonucleotides of the invention are selected from the group consisting

in the treatment of a variety of disorders, including sclerotic disorders and restenosis, associated with the inappropriate synthesis of extracellular matrix proteins, particularly collagen.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 36 OF 41 USPATFULL

2001:197035 USPATFULL ACCESSION NUMBER: Tyrosine kinase inhibitors TITLE:

INVENTOR(S):

Fraley, Mark E., North Wales, PA, United States Hartman, George D., Lansdale, PA, United States Hartman, Randall W., Newbury Park, CA, United States Merck & Co., Inc., Rahway, NJ, United States (U.S.

PATENT ASSIGNEE(S): corporation)

NUMBER KIND DATE 20011106 PATENT INFORMATION: us 6313138 R1

US 2001047007 20011129 Α1 US 2001-788720 20010220 APPLICATION INFO.: (9)

> NUMBER DATE

PRIORITY INFORMATION: US 2000-185024P 20000225 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED

PRIMARY EXAMINER: Dentz, Bernard

LEGAL REPRESENTATIVE: Garcia-Rivas, J. Antonio, Daniel, Mark R.

NUMBER OF CLAIMS: 32 EXEMPLARY CLAIM: 1 LINE COUNT: 2167

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to compounds which inhibit, regulate and/or modulate tyrosine kinase signal transduction, compositions which contain these compounds, and methods of using them to treat tyrosine kinase-dependent diseases and conditions, such as angiogenesis, cancer, tumor growth, atherosclerosis, age related macular degeneration, diabetic retinopathy, inflammatory diseases, and the like in mammals.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 37 OF 41 USPATFULL

ACCESSION NUMBER: 2001:185309 USPATFULL TITLE: Tyrosine kinase inhibitors

INVENTOR(S): Fraley, Mark E., North Wales, PA, United States

Arrington, Kenneth L., Elkins Park, PA, United States Bilodeau, Mark T., Lansdale, PA, United States Hartman, George D., Lansdale, PA, United States

Hoffman, William F., Lansdalé, PÁ, United States Kim, Yuntae, Harleysville, PA, United States

Hungate, Randall W., Newbury Park, CA, United States Merck & Co., Inc., Rahway, NJ, United States (U.S.

PATENT ASSIGNEE(S): corporation)

NUMBER KIND DATE PATENT INFORMATION: US 6306874 в1 20011023 us 2000-690598 APPLICATION INFO.: 20001017 (9)

NUMBER DATE

PRIORITY INFORMATION: US 1999-160356P 19991019 (60) DOCUMENT TYPE: Utility

FILE SEGMENT: GRANTED PRIMARY EXAMINER: Shah, Mukund J.

ASSISTANT EXAMINER: Truong, Tamthom N.

LEGAL REPRESENTATIVE: Garcia-Rivas, J. Antonio, Daniel, Mark R. NUMBER OF CLAIMS: 36

EXEMPLARY CLAIM: 1 LINE COUNT: 3068

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to compounds which inhibit, regulate and/or modulate tyrosine kinase signal transduction, compositions which contain these compounds, and methods of using them to treat tyrosine kinase-dependent diseases and conditions, such as angiogenesis, cancer, tumor growth, atherosclerosis, age related macular degeneration,

L11 ANSWER 38 OF 41 USPATFULL

2001:78947 USPATFULL ACCESSION NUMBER:

Antisense oligonucleotide modulation of human mdm2 TITLE:

expression

INVENTOR(S): Miraglia, Loren J., Encinitas, CA, United States

Nero, Pamela, Oceanside, CA, United States Graham, Mark J., San Clemente, CA, United States Monia, Brett P., La Costa, CA, United States

PATENT ASSIGNEE(S): Isis Pharmaceuticals, Inc., Carlsbad, CA, United States

(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 6238921 в1 20010529 US 1998-48810 APPLICATION INFO.: 19980326 (9)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Schwartzman, Robert A.

ASSISTANT EXAMINER:

Shibuya, Mark L. Law Offices of Jane Massey Licata LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 8 EXEMPLARY CLAIM: 1 LINE COUNT: 1117

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Compounds, compositions and methods are provided for inhibiting the expression of human mdm2. The compositions comprise antisense oligonucleotides targeted to nucleic acids encoding mdm2. Methods of using these oligonucleotides for inhibition of mdm2 expression and for treatment of diseases such as cancers associated with overexpression of mdm2 are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 39 OF 41 USPATFULL

ACCESSION NUMBER: 2001:75543 USPATFULL

Glucocorticoid receptor agonist and decreased PP5 TITLE: Honkanen, Richard E., Mobile, AL, United States South Alabama Medical Science Foundation, Mobile, AL, INVENTOR(S): PATENT ASSIGNEE(S):

United States (U.S. corporation)

KIND NUMBER DATE us 6235891 us 1999-282736 Utility PATENT INFORMATION: 20010522 В1 APPLICATION INFO.: 19990331 (9)

DOCUMENT TYPE: FILE SEGMENT: Granted PRIMARY EXAMINER: Yucel, Remy Schmidt, M ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Braman & Rogalskyj, LLP

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 13 Drawing Figure(s); 5 Drawing Page(s)

LINE COUNT: 1893

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A composition comprises a glucocorticoid receptor agonist and a compound which decreases levels of active human serine/threonine protein phosphatase 5 protein in cells. The glucocorticoid receptor agonist is dexamethasone and the compound is an antisense oligonucleotide of about 8 to 50 nucleotides in length which is targeted to a nucleic acid encoding human serine/threonine protein phosphatase 5. The composition is useful in a method of enhancing glucocorticoid activity, and in a method of enhancing the inhibition of hyperproliferation of cells where the inhibition is by contacting the cells with a compound which decreases levels of active human serine/threonine protein phosphatase 5 protein in cells. The compound is thus useful to enhance glucocorticoid therapy and to enhance inhibition of hyperproliferation relating to PP5.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 40 OF 41 USPATFULL

ACCESSION NUMBER: 2001:18459 USPATFULL

TITLE: Antisense modulation of human mdm2 expression INVENTOR(S): Miraglia, Loren J., Encinitas, CA, United States Graham, Mark J., San Clemente, CA, United States Monia, Brett P., La Costa, CA, United States

Cowsert, Lex M., Carlsbad, CA, United States

PATENT ASSIGNEE(S): Isis Pharmaceuticals Inc., Carlsbad, CA, United States

(U.S. corporation)

NUMBER KIND DATE PATENT INFORMATION: US 6184212 20010206 в1 us 1999-280805 19990326 (9)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1998-48810, filed

on 26 Mar 1998

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

APPLICATION INFO.:

PRIMARY EXAMINER: Elliott, George C.

ASSISTANT EXAMINER:

Epps, Janet Law Offices of Jane Massey Licata LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 20 2192 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Compounds, compositions and methods are provided for inhibiting the expression of human mdm2. The compositions include antisense compounds targeted to nucleic acids encoding mdm2. Methods of using these oligonucleotides for inhibition of mdm2 expression and for treatment of diseases such as cancers associated with overexpression of mdm2 are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L11 ANSWER 41 OF 41 CAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2000:171770 CAPLUS DUPLICATE 5

132:329861 DOCUMENT NUMBER:

TITLE: c-Myc antisense limits rat liver regeneration and

indicates role for c-Myc in regulating cytochrome

P-450 3A activity

AUTHOR(S):

Arora, Vikram; Knapp, Derek C.; Smith, Barbara L.; Statdfield, Mary L.; Stein, David A.; Reddy, Muralimohan T.; Weller, Dwight D.; Iversen, Patrick L. AVI BioPharma, Corvallis, OR, USA

CORPORATE SOURCE:

Journal of Pharmacology and Experimental Therapeutics SOURCE:

(2000), 292(3), 921-928 CODEN: JPETAB; ISSN: 0022-3565

PUBLISHER: American Society for Pharmacology and Experimental

Therapeutics

DOCUMENT TYPE: Journal LANGUAGE: English

Expression of c-myc protein is assocd. with cell proliferation. The ***antisense*** oligomers to inhibit c-myc present study uses expression in the regenerating rat liver after 70% partial hepatectomy (PH). ***Antisense*** phosphorodiamidate ***morpholino*** oligomers (novel DNA analogs) were administered i.p. immediately after surgery to block expression of c-myc within the first 24 h after PH. 20-mer PMO complimentary to the c-myc mRNA at the translation start site was an effective sequence (AVI-4126, 5'-ACGTTGAGGGGCATCGTCGC-3'). A single i.p. dose of 0.5 mg/kg AVI-4126 caused redn. of the regenerating liver c-myc protein in a sequence-specific and dose-dependent and compliment in a sequence-specific and dose-dependent. Inhibition of c-myc expression resulted in redn. of proliferating cell nuclear antigen and arrested cells in the GO/G1 phase of the cell cycle. The ratio of G2:G0 cell populations in the regenerating liver 24 h after PH dropped from 29.1 in saline vehicle-treated rats to 18.0 in rats treated with 2.5 mg/kg AVI-4126. The expression of cell cycle checkpoint protein ***p53*** was inhibited with increasing doses of AVI-4126, but protein ***p53*** was inhibited with increasing doses of AVI-4126, but expression of p21waf-1 was unaffected. The activity of cytochrome P 450 3A2 (CYP3A2) was evaluated by immunoblot anal. and erythromycin N-demethylation. AVI-4126 did not alter CYP3A activity in nonhepatectamized animals but showed a dose-dependent decrease in PH rats. We conclude that AVI-4126, ***antisense*** oligomer to c-myc, can We conclude that AVI-4126, ***antisense*** oligomer to c-myc, can reduce cell proliferation in the regenerating rat liver. Furthermore,

inhibition of c-myc may indirectly influence the expression of CYP3A.

ENCE COUNT: 29 THERE ARE 29 CITED REFERENCES AVAILABLE FOR THIS REFERENCE COUNT: RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

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FILE 'MEDLINE, CAPLUS, LIFESCI, EMBASE, USPATFULL, BIOSIS' ENTERED AT
      19:01:24 ON 21 JAN 2003
L1
L2
              4478 S P53 AND ANTISENSE
                78 S L1 AND SPLICE ACCEPTOR SITE
78 DUP REM L2 (0 DUPLICATES REMOVED)
L3
L4
L5
                 2 S L3 AND MORPHOLINO
               171 S L1 AND MORPHOLINO
L6
L7
               157 DUP REM L5 (14 DUPLICATES REMOVED) 37 S L6 AND PY<2001
L8
              2189 S P53 (P) ANTISENSE
Ŀ9
                66 S L8 AND MORPHOLINO
                54 S L9 NOT L7
L10
                41 DUP REM L10 (13 DUPLICATES REMOVED)
L11
=> s p53 (s) antisense
            1988 P53 (S) ANTISENSE
L12
=> d 112 1980-1988 ibib abs
L12 ANSWER 1980 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. ACCESSION NUMBER: 1992:404269 BIOSIS
DOCUMENT NUMBER:
                        BR43:60144
                        TOXICITY OF HUMAN
                                                ***P53***
                                                                 ***ANTISENSE***
TITLE:
                        OLIGONUCLEOTIDE INFUSIONS IN RHESUS MACACA.
                        SPINOLO J; BAYEVER E; IVERSEN P; JOHANSSON S; CORNISH K; PIRRUCELLO S; SMITH L; ARNESON M
AUTHOR(S):
CORPORATE SOURCE:
                        UNIV. NEBRASKA MED. CENTER, OMAHA, NEBR. 68198.
SOURCE:
                        83RD ANNUAL MEETING OF THE AMERICAN ASSOCIATION FOR CANCER
                        RESEARCH, SAN DIEGO, CALIFORNIA, USA, MAY 20-23, 1992. PROC AM ASSOC CANCER RES ANNU MEET, (1992) 33 (0), 523.
                        CODEN: PAMREA.
DOCUMENT TYPE:
                        Conference
FILE SEGMENT:
                        BR; OLD
                        English
LANGUAGE:
L12 ANSWER 1981 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.
                        1992:404268 BIOSIS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                        BR43:60143
                          ***ANTISENSE***
                                                  ***P53***
TITLE:
                                                                 OLIGODEOXYNUCLEOTIDES AS
                        POTENTIAL HUMAN ANTI-LEUKEMIC AGENTS.
AUTHOR(S):
                        BAYEVER E; HAINES K H; IVERSEN P L; SPINOLO J; KAY H D;
CORPORATE SOURCE:
                        UNIV. NEBRASKA MED. CENTER, OMAHA, NEBR. 68102.
83RD ANNUAL MEETING OF THE AMERICAN ASSOCIATION FOR CANCER
SOURCE:
                        RESEARCH, SAN DIEGO, CALIFORNIA, USA, MAY 20-23, 1992. PROC
AM ASSOC CANCER RES ANNU MEET, (1992) 33 (0), 523.
                        CODEN: PAMREA.
DOCUMENT TYPE:
                        Conference
FILE SEGMENT:
                        BR; OLD
English
LANGUAGE:
L12 ANSWER 1982 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. ACCESSION NUMBER: 1992:404267 BIOSIS
DOCUMENT NUMBER:
                        BR43:60142
TITLE:
                                            ***P53***
                                                             ***ANTISENSE***
                        SYSTEMIC HUMAN
                        OLIGONUCLEOTIDE IN RHESUS MONKEY.
AUTHOR(S):
                        IVERSEN P; CORNISH K; JOHANSSON S; FOY M; BERGOT J;
                        FREDIANI J; SMITH L; ARNESON M; BAYEVER É; SPINOLO J
CORPORATE SOURCE:
                        UNMC, OMAHA, NEBR. 68198.
SOURCE:
                        83RD ANNUAL MEETING OF THE AMERICAN ASSOCIATION FOR CANCER
                        RESEARCH, SAN DIEGO, CALIFORNIA, USA, MAY 20-23, 1992. PROC
AM ASSOC CANCER RES ANNU MEET, (1992) 33 (0), 522.
                        CODEN: PAMREA.
DOCUMENT TYPE:
                        Conference
FILE SEGMENT:
                        BR; OLD
LANGUAGE:
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L12 ANSWER 1983 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. ACCESSION NUMBER: 1992:336583 BIOSIS
DOCUMENT NUMBER:
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                          ***ANTISENSE***
TITLE:
                                                  ***P53***
                                                                RNA REDUCES THE TUMOR
                        SUPPRESSOR FUNCTION IN HUMAN LUNG CANCER CELL LINES
                                                             ***P53***
                        CARRYING WILD TYPE OR MUTATED
```

CORPORATE SOURCE: DEP. THORACIC SURGERY, UNIV. TEX. M. D. ANDERSON CANCER CENT., HOUSTON, TEX. 77030.

KEYSTONE SYMPOSIUM ON GENE TRANSFER, REPLACEMENT AND SOURCE:

AUGMENTATION, COPPER MOUNTAIN, COLORADO, USA, APRIL 3-9, 1992. J CELL_BIOCHEM SUPPL, (1992) 0 (16 PART F), 50.

CODEN: JCBSD7.

DOCUMENT TYPE: Conference BR; OLD English FILE SEGMENT: LANGUAGE:

L12 ANSWER 1984 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. ACCESSION NUMBER: 1991:524536 BIOSIS

DOCUMENT NUMBER: BA92:135996

RELEASE OF EARLY HUMAN HEMATOPOIETIC PROGENITORS FROM TITLE:

QUIESCENCE BY ANTISENSE TRANSFORMING GROWTH FACTOR BETA-1

OR RB OLIGONUCLEOTIDES.

HATZFIELD J; LI M-L; BROWN E L; SOOKDEO H; LEVESQUE J-P; O'TOOLE T; GURNEY C; CLARK S C; HATZFELD A LAB. C.N.R.S. BIOL. CELLULAIRE ET MOL. DES FACTEURS AUTHOR(S):

CORPORATE SOURCE:

CROISSANCE, I.C.I.G., HOP. PAUL-BROUSSE, 94802 VILLEJUIF

CEDEX, FRANCE

J EXP MED, (1991) 174 (4), 925-930. SOURCE:

CODEN: JEMEAV. ISSN: 0022-1007.

BA; OLD English FILE SEGMENT: LANGUAGE:

antisense We have used oligonucleotides to study the roles of transforming growth factor .beta. (TGF-.beta.) and the two antioncogenes, retinoblastoma susceptibility (Rb) and ***p53***, in the negative regulation of proliferation of early hematopoietic cells in culture. The

antisense TGF-.beta. sequence significantly enhanced the frequency of colony formation by multi-lineage, early erythroid, and granulomonocytic progenitors, but did not affect colony formation by late progenitors. Single cell culture and limiting dilution analysis indicated that autocrine TGF-.beta. is produced by a subpopulation of early progenitors. ***Antisense*** Rb but not ***antisense***

yielded similar results in releasing multipotential ***p53*** progenitors (colony-forming unit-granulocyte/erythroid/macrophage/megakary ocyte) from quiescence. Rb ***antisense*** could partially reverse the inhibitory effect of exogenous TGF-.beta. Anti-TGF-.beta. blocking antibodies, ***antisense*** TGF-.beta. or Rh oligonucleotide in similar effects. antibodies, ***antisense*** TGF-.beta., or Rb oligonucleotides all had similar effects. No additive effects were observed when these reagents were combined, suggesting a common pathway of action. Our results are consistent with the model that autocrine production of TGF-.beta. negatively regulates the cycling status of early hematopoietic progenitors through interaction with the Rb gene product.

L12 ANSWER 1985 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

1991:478184 ACCESSION NUMBER: BIOSIS

BA92:111944 DOCUMENT NUMBER:

COOPERATIVE EFFECT OF ***ANTIS
ANTISENSE - ***P53*** ***ANTISENSE*** -RB AND TITLE:

OLIGOMERS ON THE EXTENSION

OF LIFE SPAN IN HUMAN DIPLOID FIBROBLASTS TIG-1.

HARA E; TSURUI H; SHINOZAKI A; NAKADA S; ODA K
BIOLOGICAL SCI. TECHNOL., SCI. UNIV. TOKYO, CHIBA 278, JPN. CORPORATE SOURCE:

BIOCHEM BIOPHYS RES COMMUN, (1991) 179 (1), 528-534. CODEN: BBRCA9. ISSN: 0006-291X. SOURCE:

BA; OLD FILE SEGMENT: LANGUAGE: English

AUTHOR(S):

Normal human diploid fibroblasts, TIG-1, which have a replicative life span of about 62 population doublings (PD), tended to senesce after about 50 PD with a gradual decrease in sensitivity to serum. Treatment of TIG-1 cells with the ***antisense*** -Rb oligomer, which completely depleted the retinoblastoma susceptibility gene product (RB), extended life span by about 10 PD. Treatment with the ***antisense*** - ***p53*** oligomer alone had no effect; however, cotreatment with the ***antisense*** -Rb oligomer further potentiated the extension and the increased sensitivity to serum caused by the ***antisense*** -Rb oligomer alone, suggesting that ***p53*** and RB function in separate, yet complementary pathways that ***p53*** and RB function in separate, yet complementary pathways in signal transduction to senescence. The c-fos expression, which is presumed to be regulated negatively by RB, was not stimulated in partially senescent TIG-1 cells by treatment with the ***antisense*** -Rb oligomer.

L12 ANSWER 1986 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. ACCESSION NUMBER: 1990:90746 BIOSIS

DOCUMENT NUMBER: BA89:50097 MESSENGER RNA MATURATION IN MURINE ERYTHROLEUKEMIA CELLS

INDUCED TO DIFFERENTIATE.

AUTHOR(S): KHOCHBIN S; LAWRENCE J-J CORPORATE SOURCE:

LAB. DE BIOL. MOL. DU CYCLE CELLULAIRE, UNITE INSERM 309, DEP. DE RECHERCHE FONDAMENTALE, CEN-GRENOBLE, FRANCE. EMBO (EUR MOL BIOL ORGAN) J, (1989) 8 (13), 4107-4114. CODEN: EMJODG. ISSN: 0261-4189.

SOURCE:

FILE SEGMENT: BA; OLD LANGUAGE:

JAGE: English
A post-transcriptional control of gene expression was found to be responsible for a down-regulation of ***p53*** mRNA accompanying the induced differentiation of murine erythroleukemia (MEL) cells. Such a posttranscriptional control was governed by the induced synthesis of an RNA species (inRNA). In an attempt to find a potential candidate for such as function, we have localized the post-transcriptional regulation of ***p53*** mRNA in the nuclear compartment of the cells; then various ***p53*** mRNA in the nuclear compartment of the cells; then various fragments of the ***p53*** gene were used as probes for induced RNA(s) susceptible to interacting with ***p53*** pre-mRNA. This experimental approach allowed for the identification of a nuclear RNA molecule .apprx.

1.3 kb long, which was recognized specifically by a PStI-HindIII fragment located in the 5' part of the first intervening sequence of the ***p53*** gene. This RNA accumulated when cells were treated by the inducer concomitantly with high mol. wt ***p53*** mRNA precursors. However this RNA was not a maturation product of ***p53*** pre-mRNA as evidenced by its ***antisense*** orientation with respect to this as evidenced by its ***antisense*** orientation with respect to this RNA. Moreover it was markedly enriched in the poly(A)+ fraction. The complementary part of inRNA in the ***p53*** gene has been sequenced over .apprx. 1200 bp: no extensive bordlam complementary part of inRNA in the ***p53*** gene has been sequenced over .apprx. 1200 bp; no extensive homology was found in gene data banks but three restricted areas of the sequence were found homologous to a limited number of genes; they were themselves partially homologous to known repetitive sequences. Possible implication of such a sequence in the regulation of ***p53*** gene expression is discussed.

L12 ANSWER 1987 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

1989:74802 BIOSIS ACCESSION NUMBER:

DOCUMENT NUMBER: BA87:39200

TITLE: FUNCTIONAL ROLE OF BK VIRUS TUMOR ANTIGENS IN

TRANSFORMATION.

AUTHOR(S): NAKSHATRI H; PATER M M; PATER A

BASIC MED. SCI., FAC. MED., MEMORIAL UNIV. NEWFOUNDLAND, CORPORATE SOURCE:

ST. JOHN'S, NEWFOUNDLAND, CAN. A1B 3V6. J VIROL. (1988) 62 (12), 4613-4621.

J VIROL, (1988) 62 (12), 4613-4 CODEN: JOVIAM. ISSN: 0022-538X. SOURCE:

FILE SEGMENT: BA; OLD English LANGUAGE:

We have examined the role of the human papovavirus BK virus (BKV) tumor (T) antigen(s) in the maintenance of transformation and have identified the domain of T antigen essential for transformation. BKV-transformed BHK 21 and NIH 3T3 cells expressing ***antisense*** T-antigen RNA lose their ability to grow in soft agar, indicating the need for the continued expression of T antigen for the maintenance of the transformed phenotype. Experiments using translation termination linker insertion and deletion mutagenesis of BKV T antigen demonstrate that amino acids 356 to 384 are essential for transformation. Although BKV T antigen shares 100, 95, and 82% amino acid homology with that of simian virus 40 (SV40) for the nuclear localization signal, ***p53*** -binding domain, and DNA-binding domain, respectively, the transformation domains of BKV and SV40 T antigens share only 54% homology. Also, BKV T antigen lacks a substantial portion of the ATPase domain of SV40, and our results indicate the dispensability of the remaining portion for transformation by this protein. We suggest that the differences in the amino acids in the identified transformation domains together with the differences in the ATPase domains may account for the differences in the transformation. ATPase domains may account for the differences in the transformation. potentials of the two proteins.

L12 ANSWER 1988 OF 1988 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1988:374619 BIOSIS

DOCUMENT NUMBER: BA86:58529

CONSTITUTIVE EXPRESSION OF C-FOS ANTISENSE RNA BLOCKS C-FOS TITLE:

GENE INDUCTION BY INTERFERON AND BY PHORBOL ESTER AND REDUCES C-MYC EXPRESSION IN F9 EMBRYONAL CARCINOMA CELLS.

AUTHOR(S): LEVI B-Z; OZATO K

CORPORATE SOURCE: LAB. DEV. MOL. IMMUNITY, NATL. INST. CHILD HEALTH HUM.

DEV., NATL. INST. HEALTH, BETHESDA, MD. 20892, USA. GENES DEV, (1988) 2 (5), 554-566. CODEN: GEDEEP. ISSN: 0890-9369.

SOURCE:

```
LANGUAGE:
                                   English
        To address the role of c-fos proto-oncogene we constructed a plasmid that
         allows constitutive expression of RNA complementary to c-fos mRNA, and
        stably introduced this plasmid into F9 embryonal carcinoma cells. Some F9 clones expressing c-fos ***antisense*** RNA had a reduced basal level of c-fos mRNA, and were unable to induce a c-fos mRNA as well as its
        protein when stimulated with phorbol ester or with interferon (IFN).
         Nevertheless, the ability to induce major histocompatibility class I genes
         following IFN treatment was not impaired in these clones. Clones expressing c-fos ***antisense*** RNA grew as rapidly as cont
        expressing c-fos ***antisense*** RNA grew as rapidly as control F9 cells, and underwent differentiation after retinoic acid treatment.
         Unexpectedly, constitutive expression of c-myc mRNA was reduced on average by 10-fold in clones expressing c-fos ***antisense*** RNA. However,
        by 10-fold in clones expressing c-fos ***antisense*** RNA. However, expression of the ***p53*** gene and heat shock gene hsp70 was not affected in these clones, indicating the existence of a specific regulatory linkage between c-fos and c-myc genes. Cycloheximide treatment
        led to induction of a large amount of c-fos mRNA in clones expressing c-fos ***antisense*** RNA as well as in control F9 clones. The amount of c-fos ***antisense*** RNA was also increased by cycloheximide
                                                          RNA was also increased by cycloheximide c-fos ***antisense*** RNA blocks
        treatment. We postulate that c-fos ***antisense*** RNA blocks expression of the endogenous c-fos gene by accelerating the degradation of
         c-fos mRNA and that cycloheximide treatment interferes with this
        degradation.
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                                 IVERSBY SIDSEL/AU
                      1
                      2 IVERSEN/AU
0 --> IVERSEN ?/AU
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                                 IVERSEN A/AU
                      8
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                    18
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E11
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             41
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              39 DUP REM L13 (12 DUPLICATES REMOVED)
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L15
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=> d 115 ibib
L15 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2003 ACS
                           1994:208076 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                           120:208076
TITLE:
                           Cardiovascular effects of a phosphorothioate
                           oligonucleotide with sequence antisense to
                                                                          ***p53***
                           in the conscious rhesus monkey
                                                  ***Iversen,
AUTHOR(S):
                           Cornish, Kurtis G.;
                                                               Patrick*** ; Smith,
                           Larry; Arneson, Mark; Bayever, Eliel
                           Coll. Med., Univ. Nebraska, Omaha, NE, 68198-4575, USA
Pharmacology Communications (1993), 3(3), 239-47
CORPORATE SOURCE:
SOURCE:
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DOCUMENT TYPE: Journal LANGUAGE: English

=> FIL STNGUIDE
COST IN U.S. DOLLARS
SINCE FILE TOTAL
ENTRY SESSION
FULL ESTIMATED COST 227.49 227.70

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SINCE FILE TOTAL

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)

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-9.77
-9.77

FILE 'STNGUIDE' ENTERED AT 19:24:57 ON 21 JAN 2003 USE IS SUBJECT TO THE TERMS OF YOUR CUSTOMER AGREEMENT COPYRIGHT (C) 2003 AMERICAN CHEMICAL SOCIETY, JAPAN SCIENCE AND TECHNOLOGY CORPORATION, AND FACHINFORMATIONSZENTRUM KARLSRUHE

FILE CONTAINS CURRENT INFORMATION. LAST RELOADED: Jan 17, 2003 (20030117/UP).

=> d abs
YOU HAVE REQUESTED DATA FROM FILE 'CAPLUS' - CONTINUE? (Y)/N:y

ANSWER 1 OF 1 CAPLUS COPYRIGHT 2003 ACS
Antisense phosphorothioate oligonucleotides directed at ***p53*** mRNA selectively reduce renewal of human Acute Myelogenous Leukemic (AML) stem cells in culture. This would suggest that these cells could potentially be eliminated in-vivo. However, in-vitro data cannot easily be extrapolated to the intact animal. One concern relates to untoward side effects which cannot be examd. in-vitro. The objective of this study was to det. any cardiovascular alterations in the conscious primate following administration of a phosphorothioate oligonucleotide. The specific sequence, 5'-CCCTGCTCCCCCTGGCTCC-3' (designated OL(1) ***p53***), is complementary to the mRNA encoding a 53 kDa protein called ***p53***. Seven Rhesus monkeys (Macaca mulatta) were given 5 to 150 mg/kg OL(1) ***p53*** (five infusions less than 4 h and four infusions of greater than 6 days) while cardiovascular parameters were continuously monitored. Peak plasma oligonucleotide concns. were 0.37 to 0.77 .mu.M (.beta.-elimination phase) for short infusions, and 1.31 to 4.39 .mu.M for long infusions. The monkeys tolerated the treatment well and did not show any cardiovascular effects either during or after the infusion of OL(1) ***p53*** to the conscious primate does not produce alterations in cardiovascular function.

=> e hudziak ?/au
'AU' IS NOT A VALID EXPAND FIELD CODE FOR FILE 'STNGUIDE'
The indicated field code is not available for EXPAND in this
file. To see a list of valid EXPAND field codes, enter HELP
SFIELDS at an arrow prompt (=>).

=> b medline caplus lifesci embase uspatfull biosis
COST IN U.S. DOLLARS

FULL ESTIMATED COST

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)

SINCE FILE

TOTAL

SESSION
229.70

TOTAL

ENTRY SESSION
CA SUBSCRIBER PRICE 0.00 -10.42

FILE 'MEDLINE' ENTERED AT 19:27:06 ON 21 JAN 2003

FILE 'CAPLUS' ENTERED AT 19:27:06 ON 21 JAN 2003 USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT. PLEASE SEE "HELP USAGETERMS" FOR DETAILS. COPYRIGHT (C) 2003 AMERICAN CHEMICAL SOCIETY (ACS)

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FILE 'BIOSIS' ENTERED AT 19:27:06 ON 21 JAN 2003
COPYRIGHT (C) 2003 BIOLOGICAL ABSTRACTS INC.(R)
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E1
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                                 AK ROBERT"/AU OR "HUDZIAK ROBERT M"/AU OR "HUDZIAK ROBERT MICHAE
                                 L"/AU)
=> s 116 and p53
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                             1 L16 AND P53
=> d l17 ibib abs
L17 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER:
                                                       2001:816897
                                                                                  CAPLUS
DOCUMENT NUMBER:
                                                       135:353717
TITLE:
                                                       Splice-region antisense oligonucleotide composition
                                                       and targeting the mRNA splicing
                                                       Iversen, Patrick L.;
                                                                                                        ***Hudziak, Robert***
INVENTOR(S):
PATENT ASSIGNEE(S):
                                                       Avi Biopharma, Inc., USA
                                                       PCT Int. Appl., 53 pp.
SOURCE:
                                                       CODEN: PIXXD2
DOCUMENT TYPE:
                                                       Patent
LANGUAGE:
                                                       English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
           PATENT NO.
                                                KIND DATE
                                                                                              APPLICATION NO. DATE
           WO 2001083740
                                                 Α2
                                                              20011108
                                                                                              wo 2001-us14410 20010504
                   W: AU, CA, JP, KR
RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL,
                            PT, SE, TR
PRIORITY APPLN. INFO.:
                                                                                        US 2000-202376P P 20000504
          Antisense compns. targeted against an mRNA sequence for a selected protein, at a region having its 5' end from 1 to about 25 base pairs downstream of a normal splice acceptor junction in the preprocessed mRNA, are disclosed. The antisense compd. is Rnase-inactive, and is preferably a partial of the description in the preprocessed management of the description of the preprocess of the pre
           a phosphorodiamidate-linked morpholino oligonucleotide. Such targeting is
           effective to inhibit natural mRNA splice processing, produce splice
           variant mRNAs, and inhibit normal expression of the protein.
```

=> s p53 and exon()skipping

25 P53 AND EXON(W) SKIPPING

FILE 'USPATFULL' ENTERED AT 19:27:06 ON 21 JAN 2003

CA INDEXING COPYRIGHT (C) 2003 AMERICAN CHEMICAL SOCIETY (ACS)

=> dup rem 11 PROCESSING COMPLETED FOR L1 13 DUP REM L1 (12 DUPLICATES REMOVED)

=> d 12 ibib abs tot

ANSWER 1 OF 13 **MEDLINE DUPLICATE 1**

2002072617 ACCESSION NUMBER: **MEDLINE**

DOCUMENT NUMBER:

21657315 PubMed ID: 11799138
Splicing mutations in TP53 in human squamous cell carcinoma TITLE:

lines influence immunohistochemical detection.

Eicheler Wolfgang; Zips Daniel; Dorfler Annegret; Grenman Reidar; Baumann Michael **AUTHOR:**

CORPORATE SOURCE: Department of Radiotherapy and Radiation Oncology,

University, Turku, Finland.. wolfgang.eicheler@mailbox.tu-

dresden.de

JOURNAL OF HISTOCHEMISTRY AND CYTOCHEMISTRY, (2002 Feb) 50 SOURCE:

(2) 197-204.

Journal code: 9815334. ISSN: 0022-1554.

United States PUB. COUNTRY:

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: ENTRY MONTH: Priority Journals

200203

ENTRY DATE: Entered STN: 20020125

Last Updated on STN: 20020320

Entered Medline: 20020319

The mutational status of the tumor suppressor gene TP53 is often examined by immunohistochemistry. We compared the incidence of TP53 mutations in 12 permanent squamous cell carcinoma lines of the head and neck with the immunohistochemical staining obtained with two different antibodies. The mutational status of the TP53 gene was assessed by sequencing the complete coding frame of the TP53 mRNA. All 12 tumor cell lines had TP53 mutations. Six of them showed missense mutations and five had premature stop codons

caused either by splicing mutations or nonsense mutations or by ***exon*** ***skipping*** One tumor cell line was hete . One tumor cell line was heterozygous, with a truncating splicing mutation and an additional missense mutation located on different alleles. In one case, an in-frame insertion of 23 extra codons was found. All missense mutations were positive in immunhistochemistry and western blotting. The truncated ***p53*** not immunohistochemically detected in three cases with the DO-7 antibody and in five cases with the G59-12 antibody, giving false-negative results in 25% or 40%, respectively, of all tumor cell lines examined. We conclude that splicing mutations are common in squamous cell carcinoma lines and that the incidence of ***p53*** inactiviation by erroneous splicing is higher than yet reported. Sequencing of only the exons of TP53 may miss intronic mutations leading to missplicing and may therefore systematically underestimate the TP53 mutation frequency.

ANSWER 2 OF 13 USPATFULL

ACCESSION NUMBER: 2001:116764 USPATFULL

TITLE:

INVENTOR(S):

Ataxia-telangiectasia gene and its genomic organization Shiloh, Yosef, Tel Aviv, Israel Ramot-University Authority for Applied Research and PATENT ASSIGNEE(S):

Industrial Development, Tel Aviv, Israel (non-U.S.

corporation)

NUMBER KIND DATE PATENT INFORMATION: US 6265158 В1 20010724 wo 9636691 19961121 US 1998-952014 APPLICATION INFO.: 19980202 (8) WO 1996-US7025 19960516 19980202

PCT 371 date 19980202 PCT 102(e) date

RELATED APPLN. INFO.:

Continuation-in-part of Ser. No. US 1996-629001, filed on 8 Apr 1996, now patented, Pat. No. US 58661 Continuation-in-part of Ser. No. US 1995-441822, filed

on 16 May 1995, now patented, Pat. No. US 5756288

DOCUMENT TYPE: Utility

FILE SEGMENT: GRANTED

PRIMARY EXAMINER: Jones, W. Gary ASSISTANT EXAMINER: Goldberg, Jeanine LEGAL REPRESENTATIVE: Kohn & Associates

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1,7 LINE COUNT: 3109

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A purified and isolated gene, designated ATM, mutations of which cause ataxia-telangiectasia, its genomic organization, methods for the detection of the defective gene, the purified polypeptide encoded by the defective gene, and antibodies recognizing the defective protein.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 3 OF 13 USPATFULL

ACCESSION NUMBER:

PATENT ASSIGNEE(S):

2001:48208 USPATFULL

TITLE:

Ataxia-telangiectasia gene Shiloh, Yosef, Tel Aviv, Israel

INVENTOR(S):

Tagle, Danilo A., Gaithersburg, MD, United States Collins, Francis, Rockville, MD, United States The United States of America as represented by the

Department of Health and Human Services, Washington,

DC, United States (U.S. government)
Ramot University Authority for Applied Research and
Industrial Dev., Israel (non-U.S. corporation)

NUMBER KIND DATE PATENT INFORMATION: US 6211336 В1 20010403 wo 9636695 19961121 APPLICATION INFO.: us 1998-952127 19980226 (8) wo 1996-us7040 19960516

19980226 PCT 371 date 19980226 PCT 102(e) date

RELATED APPLN. INFO.:

Continuation-in-part of Ser. No. US 1995-508836, filed on 28 Jul 1995, now patented, Pat. No. US 5777093 Continuation-in-part of Ser. No. US 1995-493092, on 21 Jun 1995, now patented, Pat. No. US 5728807 Continuation-in-part of Ser. No. US 1995-441822, filed on 16 May 1995, now patented, Pat. No. US 5756288

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

Cintins, Marianne M. Delacroix-Muirheid, C. PRIMARY EXAMINER: ASSISTANT EXAMINER: LEGAL REPRESENTATIVE: Kohn & Associates

NUMBER OF CLAIMS: EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

10 Drawing Figure(s); 4 Drawing Page(s) LINE COUNT: 2279

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

There is provided a purified amino acid sequence selected from the group of Sequence ID No.: 3 and analogs thereof and mutations of Sequence ID No.: 3 which cause ataxia-telangiectasia. Also provided is a purified amino acid sequence as set forth in Sequence ID No.: 3 and analogs thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 4 OF 13 USPATFULL

ACCESSION NUMBER:

2001:36598 USPATFULL

TITLE:

Mutated forms of the ataxia-telangiectasia gene and

method to screen for a partial A-T phenotype Shiloh, Yosef, Tel Aviv, Israel

INVENTOR(S):

PATENT ASSIGNEE(S):

Ramot-University Authority for Applied Research and Industrial Development Ltd., Tel Aviv, Israel (non-U.S.

corporation)

NUMBER KIND DATE

PATENT INFORMATION: APPLICATION INFO.:

us 6200749 В1 20010313

RELATED APPLN. INFO.:

US 1996-642274 19960503 (8) Continuation-in-part of Ser. No. US 1996-629001, filed

on 8 Apr 1996 Continuation-in-part of Ser. No. US 1995-441822, filed on 16 May 1995, now patented, Pat.

No. US 5756288

DOCUMENT TYPE: FILE SEGMENT:

Utility Granted

PRIMARY EXAMINER: LEGAL REPRESENTATIVE: Arthur, Lisa B. Kohn & Associates

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

1,4

LINE COUNT: 3090

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A purified and isolated gene, designated ATM, is described mutations of which cause ataxia-telangiectasia and its genomic organization.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

DUPLICATE 2 ANSWER 5 OF 13 MEDLINE

ACCESSION NUMBER: 2000497773 MEDLINE

PubMed ID: 10980610 DOCUMENT NUMBER: 20438181

Detection of PTEN nonsense mutation and psiPTEN expression TITLE: in central nervous system high-grade astrocytic tumors by a

yeast-based stop codon assay.

Zhang C L; Tada M; Kobayashi H; Nozaki M; Moriuchi T; Abe H Section of Neurosurgery, Department of Neuropathophysiology, Hokkaido University Graduate School **AUTHOR:**

CORPORATE SOURCE:

of Medicine, Sapporo, 060-8638 Japan. ONCOGENE, (2000 Sep 7) 19 (38) 4346-53. Journal code: 8711562. ISSN: 0950-9232. SOURCE:

ENGLAND: United Kingdom PUB. COUNTRY:

Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

English LANGUAGE:

FILE SEGMENT: Priority Journals

200010 ENTRY MONTH:

Entered STN: 20001027 ENTRY DATE:

Last Updated on STN: 20001027

Entered Medline: 20001019

we have developed a new yeast-based assay for the detection of PTEN AΒ nonsense mutation, and applied it to a total of 42 astrocytic tumors. The assay utilizes homologous recombination of PCR-amplified PTEN cDNA samples to a yeast vector which expresses an in-frame PTEN::ADE2 chimera protein. An allele of nonsense mutation in the sample PTEN mRNA gives a truncated chimera protein in a yeast cell, resulting in the formation of a red colony. The assay and subsequent sequence analysis demonstrated nonsense mutations as red colonies of more than 10% in one of 10 anaplastic astrocytomas and six of 18 glioblastomas, but none in six pilocytic astrocytomas or in eight astrocytomas. Sequence analysis of white colonies showed one missense mutation in a glioblastoma. Interestingly, four of seven nonsense mutations were frame-shifts due to ***exon***

seven nonsense mutations were frame-shifts due to ***exon***

skipping . In addition, pink colonies were found in one of six
pilocytic astrocytomas, three of eight astrocytomas, two of 10 anaplastic astrocytomas, and 10 of 18 glioblastomas. Sequence analysis of the pink colonies revealed a sequence similar to those reported as psiPTEN/PTH2. By testing mRNA and genomic DNA, it was found to be a processed pseudogene which was transcribed. The psiPTEN expression was complementary to PTEN mutation, for 14 of 18 glioblastomas showed either PTEN mutation or psiPTEN expression and only one case showed both PTEN mutation and psiPTEN expression (P<0.046), suggesting a pathological role of psiPTEN expression as an alternative to PTEN mutation in glioblastomas.

ANSWER 6 OF 13 **MEDLINE**

ACCESSION NUMBER: 2000307560 MEDLINE

DOCUMENT NUMBER: PubMed ID: 10848880 20307560

TITLE: FHIT and TSG101 in thyroid tumours: aberrant transcripts

reflect rare abnormal RNA processing events of uncertain

pathogenetic or clinical significance.

AUTHOR:

McIver B; Grebe S K; Wang L; Hay I D; Yokomizo A; Liu W; Goellner J R; Grant C S; Smith D I; Eberhardt N L Department of Medicine, Mayo Clinic/Foundation, New

CORPORATE SOURCE:

zealand.

CLINICAL ENDOCRINOLOGY, (2000 Jun) 52 (6) 749-57. Journal code: 0346653. ISSN: 0300-0664. SOURCE:

PUB. COUNTRY: ENGLAND: United Kingdom

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

Priority Journals FILE SEGMENT: ENTRY MONTH:

200008

ENTRY DATE: Entered STN: 20000811

Last Updated on STN: 20000811 Entered Medline: 20000803

OBJECTIVE: The chromosomal regions containing the two putative tumour suppressors, fragile histidine triad gene (FHIT) and tumour suppressor gene 101 (TSG101), are deleted frequently in thyroid tumours. We therefore analysed FHIT and TSG101 transcripts in a group of advanced thyroid tumours to establish their role in thyroid tumorigenesis. DESIGN: Retrospective analysis of FHIT and TSG101 mRNA transcripts and genomic DNA

level not to be mutated in this cohort of tumours, served as a control. PATIENTS: We analysed nine follicular thyroid carcinomas (FTC), six papillary thyroid carcinomas and six follicular adenomas (FA) and histologically normal thyroid tissue from four of the FA patients.

MEASUREMENTS: Single stage and nested reverse transcription polymerase chain reaction (RT-PCR) products of FHIT, TSG101, and TP53 were analysed by agarose or polyacrylamide gel electrophoresis and sequenced. Genomic DNA was also analysed by polymerase chain reaction and sequencing (FHIT) or by Southern blotting (TSG101). Clinical data were correlated with the results of the mutation analysis. RESULTS: Truncated FHIT transcripts were observed frequently alongside full length transcripts with nested RT-PCR, observed frequently alongside full length transcripts with nested RT-PCR, most often in FTC, while single stage RT-PCR revealed only normal length transcripts in all tumours. Similar results were obtained for TP53, while abnormal TSG101 transcripts were detectable by single stage RT-PCR. Sequence analysis of the truncated FHIT and TSG101 transcripts revealed mainly ***exon*** ***skipping*** and alternate RNA processing events. Only a single point mutation (of TSG101) was found. Southern blotting for the TSG101 gene, and PCR amplification and sequencing of the FHIT gene showed no evidence of genomic abnormalities in either case, and there was no evidence of splice site mutations in the FHIT gene, suggesting that the truncated transcripts result from altered RNA suggesting that the truncated transcripts result from altered RNA processing. There was no relationship between tumour stage, grade or survival and the presence of FHIT or TSG101 abnormalities. CONCLUSIONS: Truncated FHIT and TSG101 transcripts in thyroid tumours reflect alternate mRNA splicing events, rather than genomic deletions. Such abnormal RNA processing seems to be common and widespread in thyroid neoplasms, as similar results were obtained by analysis of transcripts of TP53, which we had previously shown not to be mutated in these specimens. Although a pathogenetic role for these aberrant transcripts remains possible, no correlation was found with stage, histological grade or outcome in this small group of advanced thyroid malignancies. Relaxation of mRNA splice control appears to be a feature of follicular cell-derived thyroid neoplasms.

ANSWER 7 OF 13 USPATFULL

1999:16738 USPATFULL ACCESSION NUMBER:

TITLE: Computational analysis of nucleic acid information

defines binding sites

INVENTOR(S): Schneider, Thomas D., Frederick, MD, United States

Rogan, Peter K., Lebanon, PA, United States

PATENT ASSIGNEE(S): The United States of America as represented by the

Department of Health and Human Services, Washington,

DC, United States (U.S. government)

NUMBER KIND DATE US 5867402 19990202 US 1995-494115 19950623 (8)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Ramirez, Ellis B.

ASSISTANT EXAMINER: Kemper, M.

Morgan & Finnegan, L.L.P. LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

PATENT INFORMATION: APPLICATION INFO.:

30 Drawing Figure(s); 22 Drawing Page(s)

1944 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

In accordance with the present invention, binding sites are defined based upon the individual information content of a particular site of interest. Substitutions within the binding site sequences can be analyzed to determine whether the substitution will cause a deleterious mutation or a benign polymorphism. In addition, new binding sites can be identified using individual information content. Further a computer system is described for determining and displaying individual information content of a binding site sequence.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 8 OF 13 USPATFULL

ACCESSION NUMBER: 1999:4329 USPATFULL

TITLE:

INVENTOR(S):

Ataxia-telangiectasia gene and its genomic organization Shiloh, Yosef, Tel Aviv, Israel RAMOT-University Authority for Applied Research and Industrial Development, Tel Aviv, Israel (non-U.S. PATENT ASSIGNEE(S):

corporation)

NUMBER KIND DATE

19990112 PATENT INFORMATION: US 5858661 APPLICATION INFO.: US 1996-629001 19960408

(8) Continuation-in-part of Ser. No. US 1995-441822, filed on 16 May 1995, now patented, Pat. No. US 5756288 RELATED APPLN. INFO.:

DOCUMENT TYPE: Utility Granted FILE SEGMENT:

Arthur, Lisa B. Kohn & Associates PRIMARY EXAMINER: LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 6 Drawing Figure(s); 2 Drawing Page(s)

LINE COUNT: 3461

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A purified and isolated gene, designated ATM, mutations of which cause ataxia-telangiectasia and its genomic organization is disclosed. Methods and a kit for the detection of carriers of mutations of the ATM gene are provided by analysis of nucleic acids isolated from patients including in situ hybridization, Northern blotting and reverse

transcriptase--polymerase chain reaction, Southern blotting, single

strand conformational polymorphism, restriction endonuclease fingerprinting (REF), PCR amplification and DNA-chip analysis.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 9 OF 13 USPATFULL 12

ACCESSION NUMBER: 97:56495 USPATFULL

TITLE:

Methods for diagnosing cancer, precancerous state, or susceptibility to other forms of diseases by detecting an acceleration of ***exon*** ***skipping*** in

(8)

IRF-1 mRNA

Taniguchi, Tadatsugu, Ibaraki, Japan Harada, Hisashi, Suita, Japan INVENTOR(S):

Boehringer Ingelheim International GmbH, Germany, PATENT ASSIGNEE(S):

Federal Republic of (non-U.S. corporation)

NUMBER KIND DATE US 5643729 19970701 US 1995-393997 19950224

PATENT INFORMATION: APPLICATION INFO.:

> NUMBER DATE EP 1994-102839 19940224

PRIORITY INFORMATION: DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Zitomer, Stephanie W.

ASSISTANT EXAMINER: Fredman, Jeffrey Sterne, Kessler, Goldstein & Fox P.L.L.C.

LEGAL REPRESENTATIVE: NUMBER OF CLAIMS: 16

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 8 Drawing Figure(s); 8 Drawing Page(s) LINE COUNT: 963

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention concerns a novel molecular marker useful for ΔR diagnosing hematopoietic disorders, including cancers and precancerous conditions. The invention is based on the unexpected discovery that inactivation of the IRF-1 tumor suppressor gene can occur via an altered splicing pattern of the IRF-1 primary transcript. This altered splicing pattern leads to mRNAs lacking exon 2 or exons 2 and 3. The relative amounts of full-length RNA and shortened RNA molecules are significantly different in samples obtained from patients suffering from certain cancers and precancerous conditions as compared to healthy donors.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 10 OF 13 **MEDLINE** DUPLICATE 3

ACCESSION NUMBER: 97479031 MEDLINE

DOCUMENT NUMBER: 97479031 PubMed ID: 9337692

An intronic deletion in TP53 gene causes exon 6 skipping in TITLE:

breast cancer.

Voglino G; Castello S; Silengo L; Stefanuto G; Friard O; Ferrara G; Fessia L **AUTHOR:**

CORPORATE SOURCE: Department of Clinical Pathology, Ospedale Sant' Anna.

Turin, Italy.

Journal code: 9005373. ISSN: 0959-8049.

ENGLAND: United Kingdom PUB. COUNTRY:

Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 199711

ENTRY DATE: Entered STN: 19971224

Last Updated on STN: 19971224

Entered Medline: 19971103

Six hundred and thirty primary breast cancer were screened for abnormalities in exons 5, 6, 7 and 8 of the TP53 tumour suppressor gene. Analysis of the structure of the TP53 gene from the performed with the ΑB polymerase chain reaction-single strand conformation polymorphism (PCR-SSCP) method and with direct sequencing of amplified DNA. In a breast tumour case from a postmenopausal patient, we found a deletion of 36 bp in intron 5 and no immunohistochemical staining for ***p53**** . We amplified and sequenced the cDNA region between exons 4 and 7 and showed that the deletion causes the skipping of exon 6. The resulting mRNA sequence had a frameshift that yields an inactive protein with a truncated C terminus. These results show the first example of intronic deletion causing ***exon*** ***skipping*** at the TP53 gene level.

MEDLINE ANSWER 11 OF 13 **DUPLICATE 4**

96382695 ACCESSION NUMBER: MEDLINE

DOCUMENT NUMBER: 96382695 PubMed ID: 8790559

TITLE: Alterations of the RB tumour suppressor gene in

hepatocellular carcinoma and hepatoblastoma cell lines in association with abnormal ***p53*** expression.

AUTHOR:

Farshid M; Hsia C C; Tabor E National Cancer Institute, National Institutes of Health, CORPORATE SOURCE:

Bethesda, MD 20892, USA.

JOURNAL OF VIRAL HEPATITIS, (1994) 1 (1) 45-53. Journal code: 9435672. ISSN: 1352-0504. SOURCE:

ENGLAND: United Kingdom PUB. COUNTRY:

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 199610

ENTRY DATE: Entered STN: 19961025

Last Updated on STN: 19970203
Entered Medline: 19961017

Alterations in the expression of the RB tumour suppressor gene were found by Western immunoblot in three of seven hepatocellular carcinoma and hepatoblastoma cell lines. Abnormalities were detected by single-strand conformation polymorphism (SSCP) within exons 17-21 in RB cDNA from two of these three cell lines and within exons 20-21 in the third cell line. In these three cell lines with abnormal RB expression, abnormal expression of the ***p53*** tumour suppressor gene was also found, apparently the product of a mutant gene. Thus, mutations within the RB gene (or AB product of a mutant gene. Thus, mutations within the RB gene (or splice-site mutations with ***exon*** - ***skipping***) and apparent mutations of the ***p53*** gene together may have contributed to the development of three of these tumours or to the establishment of these cell lines.

ANSWER 12 OF 13 USPATFULL

ACCESSION NUMBER: 92:65884 USPATFULL

Methods and compositions for the detection of sequences TITLE:

in selected DNA molecules

LeMaistre, Anne, Humble, TX, United States Lee, Ming-Shen, Houston, TX, United States Board of Regents, The University of Texas System, Austin, TX, United States (U.S. corporation) INVENTOR(S):

NUMBER KIND DATE PATENT INFORMATION: US 5137806 19920811 APPLICATION INFO.: US 1989-448118 19891211 (7)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER: Wax, Robert A. Escallon, Miguel ASSISTANT EXAMINER: LEGAL REPRESENTATIVE: Arnold, White & Durkee

NUMBER OF CLAIMS: 22 EXEMPLARY CLAIM:

PATENT ASSIGNEE(S):

NUMBER OF DRAWINGS: 10 Drawing Figure(s); 3 Drawing Page(s)

LINE COUNT: 1498 AB The present disclosure relates to novel procedures and primers for use in conenction with PCR or in vitro DNA sequence amplification to detect sequence variants, such as sequence modifications or mutations. The invention will have particular applicability in the detection of point or other relatively short mutations where the expected location or configuration of the mutation is known. Primers of the invention incorporate a 3' terminal nucleotide or nucleotides complementary to the sequence variance, and thereby serve to successfuly prime chain elongation only on DNA templates which include the particular variant. Exemplary mutations suitable for detection through practice of the invention include those involved in beta-thalassemia, sickle cell anemia, hemoglobin C disease, diabetes, acute intermittent porphyria, lung, breast, and colon cancers and others.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 13 OF 13 DUPLICATE 5 MEDLINE

93201632 ACCESSION NUMBER: **MEDLINE**

PubMed ID: 1295697 93201632 DOCUMENT NUMBER:

Alternatively-spliced ***p53*** mRNA in the FAA-HTC1 TITLE:

rat hepatoma cell line without the splice site mutations.

Fukuda I; Ogawa K AUTHOR:

Department of Pathology, Asahikawa Medical College, Japan. CELL STRUCTURE AND FUNCTION, (1992 Dec) 17 (6) 427-32. Journal code: 7608465. ISSN: 0386-7196. CORPORATE SOURCE:

SOURCE:

PUB. COUNTRY: Japan

Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

LANGUAGE: English

FILE SEGMENT: Priority Journals

OTHER SOURCE: GENBANK-S47136; GENBANK-S47137; GENBANK-S47164;

GENBANK-S47165; GENBANK-S47166; GENBANK-S47167; GENBANK-S47168; GENBANK-S51472; GENBANK-S57234;

GENBANK-S72771

ENTRY MONTH: 199304

Entered STN: 19930507 ENTRY DATE:

Last Updated on STN: 19970203 Entered Medline: 19930420

p53 gene has been found in a rat ΑB A novel mutation of the hepatoma cell line, FAA-HTC1. This cell line carried two kinds of abnormal ***p53*** transcripts; one lacked the exon 8 sequence, and the other had a single base substitution G to T which resulted in a new stop codon in exon 8. In the genomic DNA, this base substitution in exon 8 was present, indicating that both transcripts were transcribed from the mutated gene. No mutation was detected in its two flanking introns. In this cell line, the exon-deleted transcript seems to be generated by
skipping due to an unknown mechanism other ***exon*** due to an unknown mechanism other than splice site mutations.

=> kwic 12 KWIC IS NOT A RECOGNIZED COMMAND The previous command name entered was not recognized by the system. For a list of commands available to you in the current file, enter "HELP COMMANDS" at an arrow prompt (=>).

=> d kwic 12

12 ANSWER 12 OF 13 USPATFULL

. . cystic fibrosis and Huntington's chorea. Furthermore, cancer oncogenes such as N-ras, K-ras, H-ras, Neu, or tumor suppressor genes such as ***p53*** are known to have such mutations which contribute to the cancer development. In the case of sickle cell disease, the.

DETD disease (Grandchamp, et al.: A point mutation G to A in exon 12 of the porphobilinogen deaminase gene results in ***exon***

skipping and is responsible for acute intermittent prophyria. Nucleic acids research 17:6637-49, 1989). As in example I a primer construct can.

=> s p53 and antisense and (exon or intron) 1379 P53 AND ANTISENSE AND (EXON OR INTRON)

=> s p53 same antisense same (exon or intron) MISSING OPERATOR 'SAME (EXON' The search profile that was entered contains terms or => s p53 (p) antisense (p)(exon or intron) L4 57 P53 (P) ANTISENSE (P) (EXON OR INTRON)

=> dup rem 14

PROCESSING COMPLETED FOR L4

32 DUP REM L4 (25 DUPLICATES REMOVED)

=> d 15 ibib abs tot

ANSWER 1 OF 32 USPATFULL

2002:251087 USPATFULL ACCESSION NUMBER:

METHOD OF DETECTION OF NEOPLASTIC CELLS TITLE:

INVENTOR(S): SIDRANSKY, DAVID, BALTIMORE, MD, UNITED STATES BAYLIN, STEPHEN B., BALTIMORE, MD, UNITED STATES

PATENT ASSIGNEE(S): JOHN HOPKINS UNIVERSITY SCHOOL OF MEDICINE (U.S.

corporation)

NUMBER KIND DATE

PATENT INFORMATION: us 2002137030 20020926 Α1 APPLICATION INFO.: us 1999-225904 Α1 19990105

(9) RELATED APPLN. INFO.: Division of Ser. No. US 1995-497535, filed on 30 Jun

1995, PATENTED Utility

DOCUMENT TYPE: FILE SEGMENT: APPLICATION

LISA A. HAILE, PH.D., GRAY CARY WARE & FREIDENRICH LLP, 4365 EXECUTIVE DRIVE, SUITE 1600, SAN DIEGO, CA, LEGAL REPRESENTATIVE:

92121-2189

NUMBER OF CLAIMS: 11 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 11 Drawing Page(s)

LINE COUNT: 2249

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Methylation of p16 DNA and a resultant decrease in p16 gene expression is associated with transcriptional block and is associated with a variety of neoplasms. A method for detecting a neoplasm in a subject by detecting methylation of 5'CpG islands in p16 DNA, or detecting p16 mRNA or polypeptide levels in a sample is also provided. 51

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 2 OF 32 USPATFULL

ACCESSION NUMBER: 2002:303980 USPATFULL

TITLE: Modification of mutated P53 gene in tumors by

retroviral delivery of ribozyme A

Roth, Jack A., Houston, TX, United States Cai, De Wei, Cheltenham, PA, United States INVENTOR(S):

Mukhopadhyay, Tapas, Houston, TX, United States

Board of Regents, The University of Texas System, PATENT ASSIGNEE(S):

Austin, TX, United States (U.S. corporation)

NUMBER KIND DATE US 6482803 в1 20021119 19950901 (8)

US 1995-523030 Utility APPLICATION INFO.: DOCUMENT TYPE:

FILE SEGMENT: GRANTED PRIMARY EXAMINER:

LeGuyader, John L. LEGAL REPRESENTATIVE: Fulbright & Jaworski

NUMBER OF CLAIMS: 25 **EXEMPLARY CLAIM:** 1,4

PATENT INFORMATION:

NUMBER OF DRAWINGS: 12 Drawing Figure(s); 7 Drawing Page(s)

LINE COUNT: 2784

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention discloses expression constructs and methods for employing them that result in the modulation of abnormal oncogene and tumor suppressor genes in a novel approach to cancer prevention and therapy. In one embodiment, an expression construct expresses a ribozyme that inactivates mutant p53 and also expresses the functional p53.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 3 OF 32 USPATFULL

ACCESSION NUMBER: 2002:152389 USPATFULL

TITLE: Box-dependent Myc-interacting protein (Bin1)

Prendergast, George C., Bala Cynwyd, PA, United States Sakamuro, Daitoku, West Lafayette, IN, United States INVENTOR(S):

PATENT ASSIGNEE(S):

The Wistar Institute of Anatomy and Biology,

Philadelphia, PA, United States (U.S. corporation)

NUMBER KIND DATE 20020625 PATENT INFORMATION: US 6410238 в1 wo 9855151 19981210 19991203 US 1999-445247 (9) APPLICATION INFO.: WO 1998-US11647 19980604

19991203 PCT 371 date Continuation-in-part of Ser. No. US 1997-870126, filed on 6 Jun 1997, now patented, Pat. No. US 6048702 RELATED APPLN. INFO.:

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED PRIMARY EXAMINER: McGarry, Sean LEGAL REPRESENTATIVE: Howson and Howson

NUMBER OF CLAIMS: 10 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 21 Drawing Figure(s); 19 Drawing Page(s)

LINE COUNT: 2809

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides Bin1 genomic sequences and proteins encoded thereby. Also provided are compositions and methods utilizing these sequences and proteins in the diagnosis and treatment of cancers and hyperplastic disease states. Further provided are oligonucleotides derived from sequences encoding Bin1, as well as compositions and methods utilizing same for diagnostic and therapeutic purposes.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 4 OF 32 USPATFULL

ACCESSION NUMBER: 2002:69973 USPATFULL

p53 antisense agent and method TITLE:

INVENTOR(S):

Iversen, Patrick L., Corvallis, OR, United States AVI BioPharma, Inc., Corvallis, OR, United States (U.S. PATENT ASSIGNEE(S):

corporation)

KIND DATE NUMBER US 6365577 US 1999-426804 PATENT INFORMATION: 20020402 в1 (9) APPLICATION INFO.: 19991022

> NUMBER DATE

US 1998-105695P Utility PRIORITY INFORMATION: DOCUMENT TYPE: 19981026 (60)

FILE SEGMENT: GRANTED PRIMARY EXAMINER: Wang, Andrew ASSISTANT EXAMINER: Zara, Jane LEGAL REPRESENTATIVE: Gorthey, LeeAnn

NUMBER OF CLAIMS: 18 **EXEMPLARY CLAIM:**

2 Drawing Figure(s); 2 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 1006

CAS INDEXING IS AVAILABLE FOR THIS PATENT. AB

Antisense oligonucleotides useful for treating a disease state characterized by p53 induction, such as proliferative cell disorders, e.g. cancer, or a hypoxic state induced by an ischemic attack, such as stroke, are described. The antisense agents are preferably of the class known as "steric blocker" type oligonucleotides, including morpholino oligonucleotides, peptide nucleic acids, 2'-O-allyl or 2'-O-alkyl modified oligonucleotides, or N3'.fwdarw.P5' phosphoramidate

oligonucleotides.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 5 OF 32 DUPLICATE 1 **MEDLINE**

ACCESSION NUMBER: 2002473200 **MEDLINE**

DOCUMENT NUMBER: 22220365 PubMed ID: 12235210 DeltaNp73, a dominant-negative inhibitor of wild-type p53 TITLE:

AUTHOR:

and TAp73, is up-regulated in human tumors.
Zaika Alex I; Slade Neda; Erster Susan H; Sansome
Christine; Joseph Troy W; Pearl Michael; Chalas Eva; Moll

Ute M

SOURCE:

Brook, NY 11794, USA.
JOURNAL OF EXPERIMENTAL MEDICINE, (2002 Sep 16) 196 (6)

765-80.

Journal code: 2985109R. ISSN: 0022-1007.

PUB. COUNTRY:

United States
Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH: ENTRY DATE:

200210

Entered STN: 20020918 Last Updated on STN: 20021011 Entered Medline: 20021010

p73 has significant homology to ***p53*** . However, tumor-associated AΒ up-regulation of p73 and genetic data from human tumors and p73-deficient mice exclude a classical Knudson-type tumor suppressor role. We report that the human TP73 gene generates an NH(2) terminally truncated isoform. DeltaNp73 derives from an alternative promoter in _***intron*** 3 and DeltaNp73 derives from an alternative promoter in ***intron*** lacks the transactivation domain of full-length TAp73. DeltaNp73 is frequently overexpressed in a variety of human cancers, but not in normal tissues. DeltaNp73 acts as a potent transdominant inhibitor of wild-type ***p53*** and transactivation-competent TAp73. DeltaNp73 efficiently counteracts transactivation function, apoptosis, and growth suppression mediated by wild-type ***p53*** and TAp73, and confers drug resistance to wild-type ***p53*** harboring tumor cells. Conversely, down-regulation of endogenous DeltaNp73 levels by ***antisense*** ***p53*** - and methods alleviates its suppressive action and enhances ***p53 TAp73-mediated apoptosis. DeltaNp73 is complexed with wild-type ***p53*** , as demonstrated by coimmunoprecipitation from cultured cells and primary tumors. Thus, DeltaNp73 mediates a novel inactivation mechanism of ***p53*** and TAp73 via a dominant-negative family network. Deregulated expression of DeltaNp73 can bestow oncogenic activity upon the TP73 gene by functionally inactivating the suppressor action of ***p53*** and TAp73. This trait might be selected for in human cancers.

ANSWER 6 OF 32 USPATFULL

ACCESSION NUMBER: 2001:182581 USPATFULL

TITLE: INVENTOR(S):

Methods for delivering compounds into a cell Unger, Evan C., Tucson, AZ, United States McCreery, Thomas, Tucson, AZ, United States

PATENT ASSIGNEE(S):

Imarx Pharmaceutical Corporation (U.S. corporation)

NUMBER KIND DATE ---- ----- ---

PATENT INFORMATION: APPLICATION INFO.:

US 2001031740 A1 20011018 US 2000-742938 A1 20001221

RELATED APPLN. INFO.:

Division of Ser. No. US 1997-841169, filed on 29 Apr 1997, PENDING Continuation-in-part of Ser. No. US 1997-785661, filed on 17 Jan 1997, ABANDONED Continuation-in-part of Ser. No. US 1996-640554, filed

on 1 May 1996, ABANDONED

DOCUMENT TYPE: FILE SEGMENT:

Utility **APPLICATION**

LEGAL REPRESENTATIVE:

Woodcock Washburn Kurtz, MacKiewicz & Norris LLP, One Liberty Place - 46th Floor, Philadelphia, PA, 19103

NUMBER OF CLAIMS:

104

EXEMPLARY CLAIM: NUMBER OF DRAWINGS: LINE COUNT:

7 Drawing Page(s) 2971

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention is directed, inter alia, to a method for delivering a compound into a cell comprising administering to the cell the compound to be delivered, an organic halide, and/or a carrier. Ultrasound may also be applied, if desired.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 7 OF 32 USPATFULL

ACCESSION NUMBER:

2001:231267 USPATFULL

TITLE:

INVENTOR(S):

Promoter smooth muscle cell expression
Parmacek, Michael S., Bryn Mawr, PA, United States
Solway, Julian, Glencoe, IL, United States

PATENT ASSIGNEE(S):

Arch Development Corporation, Chicago, IL, United

States (U.S. corporation)

NUMBER DATE KIND

US 1999-431349 19991101 (9) APPLICATION INFO.:

Division of Ser. No. US 1996-726807, filed on 7 Oct RELATED APPLN. INFO.:

1996

NUMBER DATE

PRIORITY INFORMATION: US 1995-4868P 19951005 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: **GRANTED**

PRIMARY EXAMINER:

McKelvey, Terry Fulbright & Jaworski, LLP LEGAL REPRESENTATIVE:

75 NUMBER OF CLAIMS:

EXEMPLARY CLAIM:

27 Drawing Figure(s); 21 Drawing Page(s) NUMBER OF DRAWINGS:

3926 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Disclosed is a smooth muscle cell specific promoter, the SM22.alpha. gene promoter as well as the murine cDNA and genomic SM22.alpha. nucleic acid sequences. Also disclosed are methods of preventing restenosis following balloon angioplasty and methods of treating asthma based on inhibition of smooth muscle cell proliferation by expressing cell cycle control genes, or contraction inhibiting peptides in smooth muscle

cells, under the control of the SM22.alpha. promoter.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 8 OF 32 USPATFULL L5

ACCESSION NUMBER: 2001:215223 USPATFULL

Transgenic mouse models for human bladder cancer TITLE:

INVENTOR(S): Wu, Xue-Ru, Woodside, NY, United States

Sun, Tung-Tien, Scarsdale, NY, United States New York University, New York, NY, United States (U.S. PATENT ASSIGNEE(S):

corporation)

NUMBER KIND DATE PATENT INFORMATION: us 6323390 20011127 в1 US 1998-83541 APPLICATION INFO.: 19980522 (9)

Continuation-in-part of Ser. No. US 1997-969315, filed RELATED APPLN. INFO.:

on 13 Nov 1997 Continuation-in-part of Ser. No. US 1997-969315, fill on 13 Nov 1997 Continuation-in-part of Ser. No. US 1997-907800, filed on 8 Aug 1997, now patented, Pat. No. US 6001646 Continuation-in-part of Ser. No. US 1995-464961, filed on 5 Jun 1995, now patented, Pat. No. US 5824543
Utility

DOCUMENT TYPE: GRANTED FILE SEGMENT:

PRIMARY EXAMINER: Martin, Jill D. Browdy & Neimark LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 27 Drawing Figure(s); 16 Drawing Page(s)

LINE COUNT: 1344

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A transgenic mouse, containing an oncogene or a tumor suppressor gene operably linked to a urothelium-specific promoter in its germ cells and somatic cells serves as an animal model system for human bladder cancer.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 9 OF 32 USPATFULL

INVENTOR(S):

ACCESSION NUMBER: 2001:168105 USPATFULL

TITLE: Method for promoting angiogenesis with a nucleic acid

construct comprising an SM22.alpha.0 promoter Parmacek, Michael S., Chicago, IL, United States Solway, Julian, Glencoe, IL, United States

Arch Development Corporation, Chicago, IL, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE US 6297221 US 1999-225670 PATENT INFORMATION: 20011002 в1 APPLICATION INFO.: 19990105

US 1999-225670 19990105 (9) Division of Ser. No. US 1996-726807, filed on 7 Oct RELATED APPLN. INFO.:

1996, now patented, Pat. No. US 6090618

DOCUMENT TYPE: Utility

FILE SEGMENT: GRANTED

Fulbright & Jaworski, LLP LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 15 EXEMPLARY CLAIM:

27 Drawing Figure(s); 21 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT:

3718

CAS INDEXING IS AVAILABLE FOR THIS PATENT. Disclosed is a smooth muscle cell specific promoter, the SM22.alpha. gene promoter as well as the murine cDNA and genomic SM22.alpha. nucleic acid sequences. Also disclosed are methods of preventing restenosis following balloon angioplasty and methods of treating asthma based on inhibition of smooth muscle cell proliferation by expressing cell cycle control genes, or control of the sm22 alpha promoter

cells, under the control of the SM22.alpha. promoter.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 10 OF 32 USPATFULL

ACCESSION NUMBER:

2001:158043 USPATFULL

TITLE: INVENTOR(S): Promoter for smooth muscle cell expression Parmacek, Michael S., Chicago, IL, United States

Solway, Julian, Glencoe, IL, United States

PATENT ASSIGNEE(S):

Arch Development Corporation, Chicago, IL, United

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: APPLICATION INFO.:

US 6291211 US 1999-431414 в1 20010918 19991101

RELATED APPLN. INFO.:

Division of Ser. No. US 1996-726807, filed on 7 Oct

1996, now patented, Pat. No. US 6090618

NUMBER DATE

PRIORITY INFORMATION:

US 1995-4868P

19951005 (60)

DOCUMENT TYPE: FILE SEGMENT:

Utility GRANTED

PRIMARY EXAMINER:

LEGAL REPRESENTATIVE:

McKelvey, Terry Fulbright & Jaworkski, LLP

NUMBER OF CLAIMS:

32

EXEMPLARY CLAIM: NUMBER OF DRAWINGS:

27 Drawing Figure(s); 21 Drawing Page(s)

LINE COUNT:

3788

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ΑB

Disclosed is a smooth muscle cell specific promoter, the SM22.alpha. gene promoter as well as the murine cDNA and genomic SM22.alpha. nucleic acid sequences. Also disclosed are methods of preventing restenosis following balloon angioplasty and methods of treating asthma based on inhibition of smooth muscle cell proliferation by expressing cell cycle control genes, or contraction inhibiting peptides in smooth muscle cells, under the control of the SM22.alpha. promoter.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 11 OF 32 USPATFULL

ACCESSION NUMBER:

2001:152683 USPATFULL

TITLE:

Early detection of ovarian carcinoma using p16 gene

INVENTOR(S):

PATENT ASSIGNEE(S):

O'Brien, Timothy J., Little Rock, AR, United States Shiqemasa, Kazushi, Hiroshima, Japan Board of Trustees of the University of Arkansas, Little

Rock, AR, United States (U.S. corporation)

KIND DATE NUMBER

PATENT INFORMATION: APPLICATION INFO.:

US 6287775 20010911 В1 US 1999-346200 19990701 (9)

RELATED APPLN. INFO.:

Division of Ser. No. US 1997-819358, filed on 17 Mar 1997 Continuation of Ser. No. US 1996-621180, filed on 21 Mar 1996

NUMBER DATE

PRIORITY INFORMATION:

19960321 (60)

DOCUMENT TYPE:

US 1996-41554P Utility

FILE SEGMENT:

GRANTED

PRIMARY EXAMINER:

McKelvey, Terry

Adler, Benjamin Aaron LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

AB

NUMBER OF DRAWINGS: 7 Drawing Figure(s); 7 Drawing Page(s)

LINE COUNT:

Increased expression of the p16 gene occurs early in the development of ovarian carcinomas. This invention detects change ovarian epithelium by measuring increases in p16 gene expression by a quantitative method that compares the levels of p16 mRNA and a control mRNA (.beta.-tubulin) in a subject to be tested against the levels of these substrates in normal subjects. A biological sample such as perituneal fluid containing mRNA derived from a subject's ovarian epithelium is taken from the subject to be tested. The mRNA is isolated from the sample, and complementary cDNA is prepared from the isolated mRNA. Using primers to p16 target sequences and to .beta.-tubulin control sequences, the cDNA is amplified. The resultant amplification products are quantitated as to p16 and beta.-tubulin gene sequences. The level p16 gene expression is assessed relative to expression levels in normal subjects. An increased level of p16 gene expression in a subject determined by this method is an indication of change in the subject's ovarian epithélium indicative of presence of a carcinoma.

ANSWER 12 OF 32 USPATFULL

ACCESSION NUMBER: 2001:147951 USPATFULL

TITLE: Method for modulating smooth muscle cell proliferation

INVENTOR(S): Parmacek, Michael S., Chicago, IL, United States

Solway, Julian, Glencoe, IL, United States Arch Development Corporation, Chicago, IL, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE US 6284743 PATENT INFORMATION: в1 20010904 US 2000-546550 20000410

APPLICATION INFO.: RELATED APPLN. INFO.:

Continuation of Ser. No. US 1999-258367, filed on 26 Feb 1999, now patented, Pat. No. US 6114311 Division of Ser. No. US 1996-726807, filed on 7 Oct 1996, now

patented, Pat. No. US 6090618

NUMBER DATE

PRIORITY INFORMATION: US 1995-4868P 19951005 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED

PRIMARY EXAMINER:

McKelvey, Terry Fulbright & Jaworski, LLP LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 23

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 14 Drawing Figure(s); 21 Drawing Page(s)

3738 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Disclosed is a smooth muscle cell specific promoter, the SM22.alpha. gene promoter as well as the murine cDNA and genomic SM22.alpha. nucleic acid sequences. Also disclosed are methods of preventing restenosis following balloon angioplasty and methods of treating asthma based on inhibition of smooth muscle cell proliferation by expressing cell cycle control genes, or contraction inhibiting peptides in smooth muscle

cells, under the control of the SM22.alpha. promoter.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ι5 **ANSWER 13 OF 32** MEDLINE DUPLICATE 2

ACCESSION NUMBER: 2001112715 **MEDLINE**

DOCUMENT NUMBER:

20576421 PubMed ID: 11013255 Cloning and characterization of the human TITLE: activity-dependent neuroprotective protein.

Zamostiano R; Pinhasov A; Gelber E; Steingart R A; Seroussi E; Giladi E; Bassan M; Wollman Y; Eyre H J; Mulley J C; **AUTHOR:**

Brenneman D E; Gozes I
Department of Clinical Biochemistry, Sackler Faculty of CORPORATE SOURCE:

Medicine, Tel Aviv University, Tel Aviv 69978, Israel.

JOURNAL OF BIOLOGICAL CHEMISTRY, (2001 Jan 5) 276 (1) SOURCE:

708-14.

Journal code: 2985121R. ISSN: 0021-9258.

PUB. COUNTRY: United States LANGUAGE: English

FILE SEGMENT: Priority Journals OTHER SOURCE: GENBANK-AF250860

ENTRY MONTH: 200102

ENTRY DATE:

Entered STN: 20010322 Last Updated on STN: 20010322 Entered Medline: 20010208

We have recently cloned the mouse activity-dependent neuroprotective AΒ protein (ADNP). Here, we disclose the cloning of human ADNP (hADNP) from a fetal brain cDNA library. Comparative sequence analysis of these two ADNP orthologs indicated 90% identity at the mRNA level. Several single nucleotide polymorphic sites were noticed. The deduced protein structure contained nine zinc fingers, a proline-rich region, a nuclear bipartite localization signal, and a homeobox domain profile, suggesting a transcription factor function. Further comparative analysis identified an ADNP paralog (33% identity and 46% similarity), indicating that these genes belong to a novel protein family with a nine-zinc finger motif genes belong to a movel protein ramily with a nine-zinc finger motified followed by a homeobox domain. The hADNP gene structure spans approximately 40 kilobases and includes five exons and four introns with alternative splicing of an untranslated second ***exon***. The hADNP gene was mapped to chromosome 20q12-13.2, a region associated with aggressive tumor growth, frequently amplified in many neoplasias, including breast bladder overion processis. including breast, bladder, ovarian, pancreatic, and colon cancers. hADNP mRNA is abundantly expressed in distinct normal tissues, and high expression levels were encountered in malignant cells. Down-regulation of ***antisense*** oligodeoxynucleotides up-regulated the tumor ***p53*** and reduced the viability of intestinal cancer suppressor cells by 90%. Thus, ADNP is implicated in maintaining cell survival, perhaps through modulation of ***p53*** .

ANSWER 14 OF 32 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 3

ACCESSION NUMBER: 2000:67424 CAPLUS

DOCUMENT NUMBER: 132:127702

TITLE: Inhibiting the growth of p53-deficient tumor cells by

administering the p53 gene

INVENTOR(S): Roth, Jack A.; Mukhopadhyay, Tapas; Tainsky, Michael

Board of Regents, the University of Texas System, USA PATENT ASSIGNEE(S): SOURCE: U.S., 37 pp., Cont.-in-part of U.S. Ser. No. 665,538,

abandoned. CODEN: USXXAM

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
us 6017524	Α	20000125	us 1992-960513	19921013
CA 2108144	AA	19920907	CA 1992-2108144	19920306
us 6436700	в1	20020820	US 1992-987235	19921207
us 6410010	В1	20020625	us 1993-145826	19931029
us 6143290	Α	20001107	US 1994-224232	19940407
us 5747469	Α	19980505	us 1994-233002	19940425
US 6069134	Α	20000530	us 1997-953290	19971017
US 6511847	В1	20030128	us 2000-668532	20000921
us 2003012770	Α1	20030116	us 2002-170240	20020611
PRIORITY APPLN. INFO.	:		US 1991-665538 B2	19910306
			US 1992-960513 A2	19921013
			US 1993-145826 A3	19931029
			US 1994-233002 A3	19940425

Disclosed are methods and compns. for the selective manipulation of gene expression through the prepn. of retroviral expression vectors for expressing ***antisense*** sequences, such as K-ras oncogene pressing ***antisense*** sequences, such as K-ras oncogene
antisense sequences, or sequences encoding a desired product, such
wild type ***p53*** sequences. Preferred retroviral vectors of the as wild type present invention incorporate the .beta.-actin promoter in a reverse orientation with respect to retroviral transcription. Preferred ***antisense*** RNA constructs of the present invention employ the use
antisense ***intron*** DNA corresponding to distinct ***intron*** regions of the gene whose expression is targeted for down-regulation. In an exemplary embodiment, a human lung cancer cell line (NCI-H460a) with a homozygous spontaneous K-ras mutation was transfected with a recombinant plasmid that synthesizes a genomic segment of K-ras in ***antisense*** orientation. Translation of the mutated K-ras mRNA was specifically inhibited, whereas expression of H-ras and

cells when expression of the mutated ras p21 protein was down-regulated by ***antisense*** RNA and cells remained viable. The growth of H460a tumors in nu/nu mice was substantially reduced by expressed K-ras

antisense RNA.

REFERENCE COUNT: 60 THERE ARE 60 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 15 OF 32 USPATFULL L5

2000:117692 USPATFULL ACCESSION NUMBER:

Method for modulating smooth muscle cell proliferation TITLE:

Parmacek, Michael S., Chicago, IL, United States INVENTOR(S):

Solway, Julian, Glencoe, IL, United States

Arch Development Corporation, Chicago, IL, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE

US 6114311 US 1999-258367 20000905 PATENT INFORMATION: APPLICATION INFO.: 19990226 (9)

Division of Ser. No. US 1996-726807, filed on 7 Oct RELATED APPLN. INFO.:

1996

DOCUMENT TYPE: Utility Granted FILE SEGMENT:

PRIMARY EXAMINER: McKelvey, Terry Arnold White & Durkee LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 18 **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 21 Drawing Figure(s); 21 Drawing Page(s)

LINE COUNT: 4109

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Disclosed is a smooth muscle cell specific promoter, the SM22.alpha. gene promoter as well as the murine cDNA and genomic SM22.alpha. nucleic acid sequences. Also disclosed are methods of preventing restenosis following balloon angioplasty and methods of treating asthma based on inhibition of smooth muscle cell proliferation by expressing cell cycle control genes, or contraction inhibiting peptides in smooth muscle

cells, under the control of the SM22.alpha. promoter.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 16 OF 32 USPATFULL

ACCESSION NUMBER: 2000:91773 USPATFULL

DNA constructs and viral vectors comprising a smooth TITLE:

muscle promoter

INVENTOR(S): Parmacek, Michael S., Chicago, IL, United States

Solway, Julian, Glencoe, IL, United States

PATENT ASSIGNEE(S): Arch Development Corporation, Chicago, IL, United

States (U.S. corporation)

NUMBER KIND DATE

US 6090618 PATENT INFORMATION: 20000718 APPLICATION INFO.: US 1996-726807 19961007 (8)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

PRIMARY EXAMINER:

McKelvey, Terry Arnold White & Durkee LEGAL REPRESENTATIVE: NUMBER OF CLAIMS: 62

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 21 Drawing Figure(s); 21 Drawing Page(s)

LINE COUNT: 4310

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Disclosed is a smooth muscle cell specific promoter, the SM22.alpha. gene promoter as well as the murine cDNA and genomic SM22.alpha. nucleic acid sequences. Also disclosed are methods of preventing restenosis following balloon angioplasty and methods of treating asthma based on inhibition of smooth muscle cell proliferation by expressing cell cycle control genes, or contraction inhibiting peptides in smooth muscle cells, under the control of the SM22.alpha. promoter.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 17 OF 32 USPATFULL 15

ACCESSION NUMBER: 2000:43942 USPATFULL

Murine and human box-dependent myc-interacting protein TITLE:

(Bin1) and uses therefor

Sakamuro, Daitoku, Philadelphia, PA, United States PATENT ASSIGNEE(S):

The Wistar Institute of Anatomy and Biology, Philadelphia, PA, United States (U.S. corporation)

NUMBER KIND DATE

us 6048702 PATENT INFORMATION: 20000411 us 1997-870126 19970606 (8) APPLICATION INFO.:

Continuation-in-part of Ser. No. US 1996-652972, filed on 24 May 1996, now patented, Pat. No. US 5723581 which is a continuation-in-part of Ser. No. US 1995-435454, filed on 5 May 1995, now patented, Pat. No. US 5605830 RELATED APPLN. INFO.:

DOCUMENT TYPE: Utility FILE SEGMENT: Granted PRIMARY EXAMINER: Guzo, David ASSISTANT EXAMINER: McGarry, Sean Howson and Howson LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 24

NUMBER OF DRAWINGS: 7 Drawing Figure(s); 21 Drawing Page(s)

LINE COUNT: 3611

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides antibodies raised against a Box-dependent myc-interacting polypeptide termed Bin1 or fragments thereof are provided. Also provided are compositions and methods utilizing these antibodies in the diagnosis and treatment of cancers and hyperplastic disease states. Further provided are oligonucleotides derived from sequences encoding Bin1, as well as compositions and methods utilizing same for diagnostic and therapeutic purposes.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 18 OF 32 USPATFULL

ACCESSION NUMBER: 2000:21739 USPATFULL

Transgenic animals_overexpressing MDM2 TITLE: Wasylyk, Bohdan, Illkirsch, France Tocque, Bruno, Courbevoie, France INVENTOR(S):

Alkhalaf, Moussa, Renne, France PATENT ASSIGNEE(S): Rhone-Poulenc Rorer SA, Antony Cedex, France (non-U.S.

corporation)

Institut National de la Sante et de la Recherche Medicale, Paris, France (non-U.S. corporation)

KIND NUMBER DATE ____ US 6028245 PATENT INFORMATION: 20000222 APPLICATION INFO.: US 1998-104497 19980625 (9)

> DATE NUMBER

PRIORITY INFORMATION: US 1997-51739P 19970703 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

Campell, Bruce R. PRIMARY EXAMINER: ASSISTANT EXAMINER: Baker, Anne-Marie

NUMBER OF CLAIMS: **EXEMPLARY CLAIM:**

NUMBER OF DRAWINGS: 32 Drawing Figure(s); 8 Drawing Page(s)

LINE COUNT: 1268

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ΑB The present invention relates to transgenic, non-human animals overexpressing a MDM2 gene. These animals model MDM2 over-expression associated with human tumors, display a major phenotype characterized by the severe skin disorder ichthyosis, and are useful for identifying compounds for the treatment of human disease. Therefore, the invention also relates to methods of using the animals for identifying compounds effective for the treatment of diseases of the skin and respiratory tract, and to the compounds themselves.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 19 OF 32 USPATFULL

ACCESSION NUMBER: 2000:4947 USPATFULL

TITLE: MDM2-specific antisense oligonucleotides INVENTOR(S): Chen, Jiandong, Metairie, LA, United States Agrawal, Sudhir, Shrewsbury, MA, United States PATENT ASSIGNEE(S): Hybridon, Inc., Cambridge, MA, United States (U.S.

corporation)

NUMBER KIND DATE

US 6013786 PATENT INFORMATION: 20000111 APPLICATION INFO.: US 1998-73567 19980506 (9)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1997-916384, filed

on 22 Aug 1997

Utility DOCUMENT TYPE: FILE SEGMENT: Granted PRIMARY EXAMINER: Degen, Nancy ASSISTANT EXAMINER: Wang, Andrew

LEGAL REPRESENTATIVE: McDonnell Boehnen Hulbert & Berghoff

NUMBER OF CLAIMS: 18

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 18 Drawing Figure(s); 21 Drawing Page(s)

LINE COUNT: 1809

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The invention provides methods to activate tumor suppressors. The invention provides methods to activate tumor suppressors. The invention further provides antisense oligonucleotides complementary to a portion of the MDM2-encoding RNA and methods for using such antisense oligonucleotides as analytical and diagnostic tools, as potentiators of transgenic animal studies and for gene therapy approaches, and as potential therapeutic agents. The invention also provides methods to augment and synergistically activate a tumor suppressor in conjunction with the use of a DNA-damage inducing agent.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 20 OF 32 USPATFULL

ACCESSION NUMBER: 1999:1434 USPATFULL

Method of detection of neoplastic cells TITLE:

INVENTOR(S): Sidransky, David, Baltimore, MD, United States Baylin, Stephen B., Baltimore, MD, United States The Johns Hopkins University School of Medicine, Baltimore, MD, United States (U.S. corporation) PATENT ASSIGNEE(S):

> NUMBER KIND DATE 19990105 US 5856094

US 1995-497535 19950630 APPLICATION INFO.: (8) Continuation-in-part of Ser. No. US 1995-439962, filed RELATED APPLN. INFO.:

on 12 May 1995, now patented, Pat. No. US 5767258

DOCUMENT TYPE: Utility

FILE SEGMENT: Granted

PRIMARY EXAMINER: Houtteman, Scott W. LEGAL REPRESENTATIVE: Fish & Richardson P.C.

NUMBER OF CLAIMS: 15 **EXEMPLARY CLAIM:**

PATENT INFORMATION:

NUMBER OF DRAWINGS: 25 Drawing Figure(s); 11 Drawing Page(s)

LINE COUNT: 2257

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Methylation of p16 DNA and a resultant decrease in p16 gene expression is associated with transcriptional block and is associated with a variety of neoplasms. A method for detecting a neoplasm in a subject by detecting methylation of 5'CpG islands in p16 DNA, or detecting p16 mRNA or polypeptide levels in a sample is also provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 21 OF 32 USPATFULL

1998:48564 USPATFULL ACCESSION NUMBER:

P53AS protein and antibody therefor TITLE:

INVENTOR(S): Kulesz-Martin, Molly F., Buffalo, NY, United States PATENT ASSIGNEE(S): Health Research, Inc., Buffalo, NY, United States (U.S.

corporation)

NUMBER KIND DATE PATENT INFORMATION: us 5747650 19980505 APPLICATION INFO.:

us 1996-644456 19960510 (8)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1993-100496, filed

on 2 Aug 1993

DOCUMENT TYPE: Utility FILE SEGMENT: Granted ASSISTANT EXAMINER: Bansal, Geetha P. Dunn, Michael L. LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 11

EXEMPLARY CLAIM: NUMBER OF DRAWINGS: 26 Drawing Figure(s); 11 Drawing Page(s)

LINE COUNT: 1580 CAS INDEXING IS AVAILABLE FOR THIS PATENT.

In accordance with the present invention, we have discovered and purified a protein designated herein as p53as, which protein is present in normal cells of a mammal and is essentially identical to known normal growth controlling protein p53 of the same mammal, at least until the final 50 amino acids of the carboxy terminal end of the protein. The invention further includes an antibody specific for protein p53as, which antibody is designated herein as Ab p53as. The antibody may be either a monoclonal or polyclonal antibody and burnay be specific for p53as of any

particular mammal such as mice and humans.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 22 OF 32 **MEDLINE DUPLICATE 4**

ACCESSION NUMBER: 1999053919 MEDLINE

PubMed ID: 9840184 DOCUMENT NUMBER: 99053919

TITLE:

The presence of wild-type TP53 is necessary for the radioprotective effect of the Bowman-Birk proteinase

inhibitor in normal fibroblasts.

Dittmann K H; Gueven N; Mayer C; Ohneseit P; Zell R; Begg A AUTHOR:

C; Rodemann H P

Départment of Radiotherapy, Eberhard-Karls-University, CORPORATE SOURCE:

Tubingen, Germany.
RADIATION RESEARCH, (1998 Dec) 150 (6) 648-55. SOURCE:

Journal code: 0401245. ISSN: 0033-7587.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals; Space Life Sciences

ENTRY MONTH: 199812

ENTRY DATE: Entered STN: 19990115

Last Updated on STN: 19990115 Entered Medline: 19981214

In the present study we have demonstrated that the Bowman-Birk proteinase inhibitor (BBI) protected normal fibroblasts from a radiation-induced reduction in cell survival, whereas in transformed fibroblasts no AB radioprotective effect was observed. It was shown that BBI reduced the radiation-induced protein stabilization and DNA-binding activity of TP53 (formerly known as ***p53***) in normal fibroblasts. In transformed fibroblasts, BBI failed to induce these effects. The analysis of the TP53 gene in transformed fibroblasts revealed a mutation in ***exon*** 5.

As a consequence of this mutation, the expression of the TP53 downstream gene CDKN1A (p21/WAF1/Cip1) is blocked. Based on experiments using TP53
antisense oligonucleotides, the radioprotective effect of BBI
could be correlated with the function of wild-type TP53. Thus BBI can be considered as a selective radioprotective agent for normal human fibroblasts.

ANSWER 23 OF 32 CAPLUS COPYRIGHT 2003 ACS SSION NUMBER: 1997:216970 CAPLUS **DUPLICATE 5**

ACCESSION NUMBER:

DOCUMENT NUMBER: 126:288623

AUTHOR(S):

TITLE: A sensitive and high-resolution method for detection

of mutations in the p53 gene using multiple fluorescence-based symmetric PCR-SSCP analysis Katsuragi, Kiyonori; Chiba, Wataru; Ikeda, Sadao; Ueta, Chie; Kinoshita, Moritoshi Diagnostics Division, Otsuka Pharmaceutical Co., Ltd., Tokushima, 771-01, Japan Biomedical Research (1997), 18(1), 57-64 CODEN: BRESD5; ISSN: 0388-6107 Riomedical Research Foundation

CORPORATE SOURCE:

SOURCE:

PUBLISHER: Biomedical Research Foundation

DOCUMENT TYPE: Journal English LANGUAGE:

p53 ΑB Mutations of the gene are an important feature of neoplastic progression in humans. The presence of such mutations has been detected in exons 5 through 8, which contain 86% of all mutations reported for the ***p53*** gene. The authors have developed a simple and rapid method, multiple fluorescence-based sym. polymerase chain reaction in a single tube and single-strand conformation polymorphism anal. in one lane (MF-SPCR-SSCP) for detection of mutations in exons 5, 6, 7 and 8 of the

a single tube using mixed four-color fluorescence-labeled sense and primers. This technique enabled labeling of each of the product with a unique fluorescent. The MF-SPCR ***exon*** products were heat-denatured and applied to 7% polyacrylamide gel contg. 5% glycerol set on an automated DNA sequencer with a gel temp.-controlling system. The authors analyzed 18 specimens of lung cancer tissue with mutations in exons 5 through 7 using MF-SPCR-SSCP method. These mutations were detected even with use of only one PCR and one set of conditions for electrophoresis.

ANSWER 24 OF 32 USPATFULL

96:120741 USPATFULL ACCESSION NUMBER:

TITLE: Methods and compositions for detecting base pair

mismatches

INVENTOR(S):

Winkler, Matthew, Austin, TX, United States Goldrick, Marianna M., Pflugerville, TX, United States Ambion, Inc., Austin, TX, United States (U.S.

PATENT ASSIGNEE(S):

corporation)

NUMBER KIND DATE

US 5589329 US 1993-155937 PATENT INFORMATION: 19961231 APPLICATION INFO.: 19931115 (8)

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

Jones, W. Gary Marschel, Ardin H. Arnold White & Durkee PRIMARY EXAMINER: ASSISTANT EXAMINER: LEGAL REPRESENTATIVE:

NUMBER OF CLAIMS: 44 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 11 Drawing Figure(s); 11 Drawing Page(s)

LINE COUNT: 1735

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention discloses improved compositions and methods for AΒ detecting mutations, including single base changes, in nucleic acid sequences using RNase protection assays. The improvements include concomitant, dramatic reductions in the salt and RNase enzyme concentrations in the RNase digestion reaction mixture which result in greater sensitivity in detecting genetic mutations. Another embodiment of the present invention is kits to be used for the detection of single base mismatches in nucleic acid samples.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 25 OF 32 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V.DUPLICATE 6

95354258 ACCESSION NUMBER: **EMBASE**

1995354258 DOCUMENT NUMBER:

TITLE: Role of p53 in MCF-10F cell immortalization and

AUTHOR: CORPORATE SOURCE:

chemically-induced neoplastic transformation.
Barnabas N.; Moraes R.; Calaf G.; Estrada S.; Russo J.
Breast Cancer Research Laboratory, Fox Chase Cancer Center,
7701 Burholme Avenue, Philadelphia, PA 19111, United States
International Journal of Oncology, (1995) 7/6 (1289-1296). SOURCE:

ISSN: 1019-6439 CODEN: IJONES

COUNTRY: Greece

DOCUMENT TYPE: Journal: Article FILE SEGMENT: 016 Cancer

LANGUAGE: English **SUMMARY LANGUAGE:** English

The present study was undertaken to determine the role of the tumor suppressor gene ***p53*** in the transformation of the human breast epithelial cell (HBEC) line MCF-10F treated with chemical carcinogens in vitro. MCF-10F is a spontaneously immortalized diploid HBEC line, derived from a mortal cell strain designated MCF-10M. MCF-10F cells became neoplastically transformed by in vitro treatment with the chemical carcinogens 7, 12-dimethylbenz(a)anthracene (DMBA) and benzo(a)pyrene (BP). DMBA and BP-treated cells gave rise to clones D3, D3-1, BP1 and BP1-E, respectively, all of which expressed colony formation in agar-methocel and high chemoinvasion index. BP1-E cells, derived from BP1, were tumorigenic in severe combined immunodeficient (SCID) mice. were tumorigenic in severe combined immunodeficient (SCID) mice. We designed this work utilizing this model in which isolated clones of cells express different stages of progression to neoplastic transformation for determining whether any specific phenotype was associated with alteration in the ***p53*** tumor supressor gene. For this purpose, Southern blot, Northern blot, single-strand conformation polymorphism (SSCP) and DNA sequencing were used to detect mutations in the highly conserved exons the cells tested by Southern and Northern blot, SSCP analysis showed a conformational shift in ***exon*** 7 in the MCF-10F cell line, and in clones BP1, BP1-E, D3, and D3-1, derived from DMBA and BP treated cells, respectively. This shift was absent in MCF-10M cells, the mortal cells from which the MCF-10F immortal cells were derived, and in the placental DNA used as control. Sequence analysis using asymmetric PCR-amplified products of ***exon*** 7 and an ***antisense*** primer revealed an insertional mutation of thymine at codon 254 in MCF-10F cells and in transformed cells, but not in MCF-10M. These data indicate that the emergence of the immortalized phenotype was associated with a mutation of ***p53*** . DMBA- or BP-treatment did not induce additional changes in the ***p53*** gene. The fact that the precursor of the immortalized MCF-10F did not present changes in ***p53*** , may indicate that the alteration of this tumor suppressor gene could be associated with the process of cell immortalization; this, in turn, might facilitate the neoplastic transformation of the cells by chemical carcinogens.

ANSWER 26 OF 32 MEDLINE DUPLICATE 7 1.5 95186358 ACCESSION NUMBER: **MEDLINE** DOCUMENT NUMBER: 95186358 PubMed ID: 7880719 TITLE: Antisense oligonucleotides directed against p53 have antiproliferative effects unrelated to effects on p53 expression. Barton C M; Lemoine N R **AUTHOR:** CORPORATE SOURCE: Imperial Cancer Research Fund Oncology Unit, Hammersmith Hospital, London, UK. BRITISH JOURNAL OF CANCER, (1995 Mar) 71 (3) 429-37. Journal code: 0370635. ISSN: 0007-0920. SCOTLAND: United Kingdom SOURCE: PUB. COUNTRY: DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE) LANGUAGE: English FILE SEGMENT: ENTRY MONTH: Priority Journals 199504 ENTRY DATE: Entered STN: 19950425 Last Updated on STN: 19980206 Entered Medline: 19950412

Antisense oligonucleotides targeting ***p53*** have been hailed as a potentially new technique for treating patients with cancer, and there have been encouraging reports of good patient tolerance in vivo and of antiproliferative effects in vitro. However, evidence is lacking that these oligonucleotides are acting via an ***antisense*** Entered Medline: 19950412 ΑB that these oligonucleotides are acting via an interaction to modulate | ***p53*** | expression expression. We examined a oligonucleotide, directed against ***antisense*** phosphorothioate ***exon*** 10 of the TP53 gene, and a chimaeric phosphorothioatephosphodiester oligonucleotide directed against the translation initiation codon. Both failed to specifically suppress
p53 protein production in a cell-free assay system or to have any ***p53*** effect on mutant expression by human pancreatic cancer cell lines. Antiproliferative effects were apparent, especially with the phosphorothioate ***antisense*** oligonucleotide, but this was independent of the ***p53*** status of the cells (mutant, wild-type or absent) and also occurred with the control (sense and random sed) oligonucleotides. The most dramatic antiproliferative effects were seen with the 'control' phosphorothicate oligonucleotides. These findings suggest that the antiproliferative effects of some ***antisense*** suggest that the antiproliferative effects of some ***antisense*** oligonucleotides may be unrelated to expression of the gene they have been designed to target. ANSWER 27 OF 32 **DUPLICATE 8** MEDLINE ACCESSION NUMBER: 92340559 **MEDLINE** 92340559 PubMed ID: 1378845 Characterization of an endogenous RNA transcript with

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ACCESSION NUMBER: 92340559 MEDLINE 92340559 PubMed ID: 1378845

TITLE: 92340559 PubMed ID: 1378845

Characterization of an endogenous RNA transcript with homology to the antisense strand of the human c-myc gene. Celano P; Berchtold C M; Kizer D L; Weeraratna A; Nelkin B D; Baylin S B; Casero R A Jr

CORPORATE SOURCE: Johns Hopkins Oncology Center Laboratories, Baltimore, Maryland 21231. CA 51068 (NCI)

CONTRACT NUMBER: CA47492 (NCI) CA51085 (NCI)

SOURCE: JOURNAL OF BIOLOGICAL CHEMISTRY, (1992 Jul 25) 267 (21) 15092-6. Journal code: 2985121R. ISSN: 0021-9258.
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PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

GENBANK-M63383; GENBANK-M86615; GENBANK-M86616; GENBANK-M86617; GENBANK-M86618; GENBANK-M86619; GENBANK-M91159; GENBANK-M91212; OTHER SOURCE:

GENBANK-X62322

199208 ENTRY MONTH:

Entered STN: 19920911 ENTRY DATE:

Last Updated on STN: 19970203

Entered Medline: 19920826 In addition to being regulated by a complex array of cis- and trans-acting AB

factors, c-myc protooncogene expression may be modulated by
antisense RNA transcripts. Our previous studies have

RNA transcripts. Our previous studies have determined that depletion of intracellular polyamines by alpha-difluoromethylornithine results in a marked decrease in the transcription of the human c-myc gene. Because of reports that ***antisense*** transcription occurs in the 5' and 3' regions of this gene, we used a genomic clone of the human c-myc gene to ascertain whether polyamine depletion might induce an ***antisense*** RNA transcript. These studies demonstrate that polyamine depletion of the human colon cancer cell line COLO 320 results in induction of an endogenous RNA transcript with high homology to the ***antisense*** strand of the second intervening sequence (PvuII-RsaI) of the c-myc gene. Furthermore, during such depletion, steady state levels of this transcript vary inversely to the sense direction c-myc RNA. RNase protection studies suggest that the ***antisense*** transcript may arise from a different gene locus than

the c-myc gene. To further identify the origins of this RNA, a CDNA library was generated from size-selected RNA and screened with c-myc sequences. A 438-base pair cDNA was isolated with approximately 85% homology, to a 285-base region in the second ***intron*** of the of the c-myc

gene. Computer homology analysis further reveals that a 120-base region within this cDNA also has approximately 85% homology to the ***antisense*** strands of a number of genes, including the growth-related genes, N-myc, ***p53****, and thymdinae kinase. These growth-related genes, N-myc, ***p53***, and thymidine kinase. These studies provide the initial characterization of an endogenous ***antisense*** RNA transcript which could influence cell growth by

modulating the expression of c-myc and other genes.

ANSWER 28 OF 32 **DUPLICATE 9** MEDLINE

93104490 ACCESSION NUMBER: **MEDLINE**

93104490 PubMed ID: 1361370 DOCUMENT NUMBER:

Transcription factors, translocations, and leukemia. TITLE:

Nichols J; Nimer S D **AUTHOR:**

Department of Medicine, UCLA School of Medicine 90024-1678. CORPORATE SOURCE:

CONTRACT NUMBER: DK43025 (NIDDK)

BLOOD, (1992 Dec 15) 80 (12) 2953-63. Ref: 118 Journal code: 7603509. ISSN: 0006-4971. SOURCE:

United States PUB. COUNTRY:

Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

General Review; (REVIEW)

(REVIEW, ACADEMIC) English

LANGUAGE:

FILE SEGMENT: Abridged Index Medicus Journals; Priority Journals

ENTRY MONTH: 199301

ENTRY DATE:

Entered STN: 19930212 Last Updated on STN: 19950206 Entered Medline: 19930125

AB The frequent occurrence of TF gene involvement in translocations associated with leukemia is remarkable, although not yet explained. The wide variety of TFs involved in these translocations and the different stages of cellular maturation argue against a unifying mechanism.

Recombinases, active during B-cell and T-cell development, have been implicated in gene arrangements involving TCR genes and in the SIL/SCL rearrangement, which involves two genes not normally rearranged. However, other mechanisms must clearly be active in generating these molecular abnormalities and perhaps they relate to the multistep maturation and differentiation processes and continuous cell turnover seen in hematopoietic cells. The difficulties in obtaining human solid tumor samples may make it more difficult to identify translocations involving TF genes in solid tumors. Recently, the cytogenetic analysis of solid tumors has improved and specific cytogenetic abnormalities have been associated with specific types of tumors. With advanced techniques, such as fluorescent in situ hybridization (a technique that does not depend on cell growth) and PCR, abnormalities involving TF genes will be discovered. Abnormalities of TF genes, other than translocations, have been seen in a broad variety of nonhematopoietic malignancies. The ***p53*** protein has been shown to bind DNA in a sequence-specific fashion and interact with a variety of DNA tumor virus oncoproteins. The broad range of cell

abnormalities will likely be implicated in many solid tumors. We have detailed several examples of how gene rearrangements that accompany chromosomal translocations in acute leukemia can alter the expression or activity of cellular TFs. Several translocations generate fusion RNA transcripts and fusion TF proteins with altered functional characteristics. Other translocations result in the expression of a gene not normally detectable in hematopoietic cells or alter the level of its expression, or affect the promoter usage or ***exon*** structure of expression, or affect the promoter usage or ***exon*** structure of the gene (Table 2). Studies are underway in many laboratories to characterize the biologic activity of these abnormal TFs and it remains to be proven that these molecular abnormalities are directly linked with leukemogenesis. The identification of abnormal fusion transcripts and proteins may allow specific therapies to be directed against "tumor-specific" DNA, mRNA, or protein targets. Therapeutic strategies based on ***antisense*** or ribozyme technology may be used to turn based on ***antisense*** or ribozyme technology may be used to turn off expression of these genes and inhibit leukemia cell growth.

Immunologic methods can also be used to direct therapy against the malignant cells.

ANSWER 29 OF 32 DUPLICATE 10 MEDLINE

ACCESSION NUMBER: **DOCUMENT NUMBER:**

94028468 MEDLINE

PubMed ID: 1340159 94028468

TITLE:

Antisense RNA and p53 regulation in induced murine cell

differentiation.

AUTHOR:

Khochbin S; Brocard M P; Grunwald D; Lawrence J J

CORPORATE SOURCE:

Laboratoire de Biologie Moleculaire du Cycle Cellulaire Unite INSERM 309, Centre d'Etudes Nucleaires de Grenoble,

SOURCE:

ANNALS OF THE NEW YORK ACADEMY OF SCIENCES, (1992 Oct 28) 660 77-87.

Journal code: 7506858. ISSN: 0077-8923.

PUB. COUNTRY:

United States

DOCUMENT TYPE:

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: FILE SEGMENT: English Priority Journals

ENTRY MONTH:

199310

ENTRY DATE:

Entered STN: 19940117 Last Updated on STN: 19970203 Entered Medline: 19931022

p53 expression is strongly modulated during the process of induced differentiation, at the same time as both cell cycle and genetic AB expression become modulated, giving rise to a commitment to terminal differentiation. We took advantage of two murine cell lines inducible for differentiation, an erythroleukemia and a melanoma cell line, to outline common features of the regulation of ***p53*** expression during the differentiation process. We found that ***p53*** mRNA decreased early after induced differentiation and that regulation was controlled at a mRNA decreased early posttranscriptional level. Our data showed that this regulation affects ***p53*** pre-mRNA maturation. Because, in both systems used, actinomycin D treatment abolished the inducer-mediated decrease of ***p53*** mRNA, we looked for induced RNAs potentially involved in this process. Using different parts of the ***p53*** gene and flanking regions as probes, we identified three RNA species whose expression is modulated during induced differentiation. A first species is made of high molecular weight RNAs that accumulate in the nuclear compartment and seem to represent ***antisense*** transcripts of the ***p53**** gene. A species 1 3-b long was found to record species whose expression is made of high molecular weight RNAs that accumulate in the nuclear compartment and seem to represent the record species are species as the record species are spec to represent ***antisense*** transcripts of the ***p53*** gene. A second species, 1.3-kb long, was found to accumulate in the nucleus of induced MEL cells and was homologous to a restricted part of the first ***intron*** of the ***p53*** gene due to the presence of a B1 repetitive element in an ***antisense*** orientation with respect to the ***p53*** pre-messenger RNA. Finally, a family of B2-containing RNAs was observed in both cytoplasmic and nuclear compartments. The variation in the amounts of sense and ***antisense*** RNAs, respectively, suggested an interesting speculative model for the maturation of B2-containing pre-messenger RNAs.

ANSWER 30 OF 32 CAPLUS COPYRIGHT 2003 ACS SSION NUMBER: 1992:52757 CAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER:

116:52757

TITLE: AUTHOR(S): BstNI/NciI polymorphism of the human p53 gene (TP53)

CORPORATE SOURCE:

Chumakov, P. M.; Jenkins, J. R. Engelhardt Inst. Mol. Biol., Moscow, 117984, USSR Nucleic Acids Research (1991), 19(24), 6969

CODEN: NARHAD; ISSN: 0305-1048

DOCUMENT TYPE:

LANGUAGE:

SOURCE:

Journal English

p53 gene that can be revealed using restriction nuclease digestion of PCR-amplified DNA segment is reported. Left-hand (sense) oligonucleotide 5'-GTTGCCCAGGGTCCCCAGGCCTCTGATTCCTCACT-3' corresponded to the region of ***intron*** five 12 bp upstream of ***exon*** 6; GGGAGGCCCTTAGCCTCGTAAGCTTCA-3' corresponded to the region of ***intron*** six 165 bp downstream of ***exon*** 6. The amplified fragment was purified by spermine poth digested with RSTNI OF Noil ***intron*** six 165 bp downstream of ***exon*** 6. The amplified fragment was purified by spermine pptn., digested with BstNI or NciI restriction nucleases and subjected to electrophoresis through 2% agarose. BstNI (CCAGG) and NciI (CCGGG) cleave different allele fragments of the Allele frequencies calcd. from 56 unrelated Caucasians were K1 = 0.31 and restriction site polymorphic A/G nucleotide responsible for BstNI/NciI downstream of ***exon*** 6 of the ***p53*** gene (17p13). Mendelian inheritance was demonstrated in 2 three-generation families.

ANSWER 31 OF 32 CAPLUS COPYRIGHT 2003 ACS SSION NUMBER: 1990:566707 CAPLUS ACCESSION NUMBER: DOCUMENT NUMBER: 113:166707 AccII polymorphism of the p53 gene
De la calle-Martin, Oscar; Fabregat, Virginia; Romero,
Matilde; Soler, Jesus; Vives, Jordi; Yague, Jordi
Servei Immunol., Hosp. Clin., Barcelona, 08036, Spain
Nucleic Acids Research (1990), 18(16), 4963
CODEN: NARHAD; ISSN: 0305-1048 TITLE: AUTHOR(S): CORPORATE SOURCE: SOURCE: DOCUMENT TYPE: LANGUAGE: English A rapid and simple method is reported to analyze an AccII polymorphism within the human ***p53*** gene using the polymerase chain reaction A rapid and simple method is reported to analyze an ACCII polymorphism within the human ***p53*** gene using the polymerase chain reaction. PCR Primers: The primer sequences corresponded to the 4th ***exon*** of the human ***p53*** gene: Sense oligo 5'-AATGGATGATTTGATGCTGTCCC-3' and ***Antisense*** oligo 5'-CGTGCAAGTCACAGACTTGGC-3'. An ACCII (CGCG) digest of the amplified fragment identifies 2 alleles; A1 = 259 bp and A2 = 160 bp + 99 bp. Allele frequencies were calcd. from 90 unrelated caucasians. A1 = 0.32 A2 = 0.68. The polymorphic AccII recognition site occurs within the 4th ***exon*** of the human ***p53*** locus

ANSWER 32 OF 32 CAPLUS COPYRIGHT 2003 ACS SSION NUMBER: 1990:71437 CAPLUS L 5 ACCESSION NUMBER: DOCUMENT NUMBER:

112:71437

TITLE:

An antisense RNA involved in p53 mRNA maturation in murine erythroleukemia cells induced to differentiate

AUTHOR(S): CORPORATE SOURCE: SOURCE:

CODEN: EMJODG; ISSN: 0261-4189

DOCUMENT TYPE: LANGUAGE:

English

A post-transcriptional control of gene expression was found to be A post-transcriptional control of gene expression was found to be responsible for a down-regulation of p53 mRNA accompanying the induced differentiation of murine erythroleukemia (MEL) cells. Such a post-transcriptional control was governed by the induced synthesis of an RNA species (inRNA). In an attempt to find a potential candidate for such a function, the post-transcriptional regulation of p53 mRNA was localized in the nuclear compartment of the cells: then various fragments of the p53 a function, the post-transcriptional regulation of postmkNA was localized in the nuclear compartment of the cells; then various fragments of the p53 gene were used as probes for induced RNA(s) susceptible to interacting with p53 pre-mRNA. This exptl. approach allowed for the identification of a nuclear RNA mol., .apprx.1.3 kb long, which was recognized specifically by a PstI-HindIII fragment located in the 5' part of the first intervening sequence of the p53 gene. This RNA accumulated when cells were treated by sequence of the p53 gene. This RNA accumulated when cells were treated by the inducer concomitantly with high mol. wt. p53 mRNA precursors. However this RNA was not a maturation product of p53 pre-mRNA as evidenced by its antisense orientation with respect to this RNA. Moreover it was markedly enriched in the poly(A) traction. The complementary part of input in the enriched in the poly(A)+ fraction. The complementary part of inRNA in the p53 gene has been sequenced over .apprx.1200 bp; no extensive homol. was found in gene data banks but 3 restricted areas of the sequence were found homologous to a limited no. of genes; they were themselves partially homologous to known B1 repetitive sequences. Possible implication of such a sequence in the regulation of p53 gene expression is discussed.

> kwic 2 4 WIC IS NOT A RECOGNIZED COMMAND he previous command name entered was not recognized by the system. => d kwic 2 4 L5 ANSWER 2 OF 32 USPATFULL In the recombinant retroviral vectors, the orientation (a) vector showed higher levels of ***p53*** expression. It is contemplated that other retroviral promoters in the construct will suppress the .beta.-actin promoter, as described in other. . . all promoters are aligned in the same direction of transcription (Emerman & Temin, 1984). Another possible explanation is that the ***intron*** and its enhancer in DETD the .beta.-actin promoter are spliced out of the retroviral message during the first round of retroviral. . . therefore this effect may have some degree of promoter specificity (Emerman & Temin, 1984; Gunning et al., 1987). If some ***antisense*** transcripts were produced in orientation (a), alternate transcripts should have been detected by Northern analysis. However, these transcripts were not detected. The effectiveness in expression of functional ***p53*** protein by the state of the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of functional ***p53*** protein by the effectiveness in expression of the effectiveness in protein by the orientation (a) construct supports the absence of inhibition by ***antisense*** . The use of .beta.-actin promoter in orientation (b) with an LNL6 retrovirus yielded low rates of infectivity and low levels of gene expression (Owens & Boyd, 1991). Therefore, to maximize expression of ***p53*** , it may be advantageous to utilize different transcriptional orientations for the genes inserted in the retroviral vector. . . . the presence of 8 .mu.g/mL polybrene. This transinfection was repeated once daily for 3 days. To examine whether the transduced ***p53*** gene was expressed in these cells; the reverse DETD gene was expressed in these cells; the reverse transcription-PCR analysis used sense primers for .beta.-actin promoter sequences 5' to the promoter/ ***p53*** junctional sequences and an opposing ***p53*** cDNA ***antisense*** primer located within ***p53*** dDNA ***antisense*** primer located within ***p53*** 4. These primers are specific for the retrovices.

primer located within ocated within ***p53*** ***p53*** CDNA ***antisense*** primer located within ***p53***

exon 4. These primers are specific for the retrovirally
transduced ***p53*** . PCR products were evaluated by Southern blot
hybridization with a .sup.32p-labeled, nick-translated ***p53***

CDNA probe. A .beta.-actin/ ***p53*** segment was detected in H226Br
cells transduced with wt- ***p53***, whereas it was not present in
parental and LNSX virus-infected cells. Western blot analysis
demonstrated detectable levels of ***p53*** protein following LNp53B
retroviral infection in ***p53*** -negative H358a cells.

An ***antisense*** ***p53*** RNA probe was synthesized as above
from a plasmid containing a ***p53*** CDNA template. Amplimers
corresponding to **exon*** 5 [5'-TACTCCCTTGCCCTCAACAAG-3' (SEQ ID
NO:25)] and **exon*** 8 [5'-CTTAGTGCTCCCTTGGGGGCAG-3' (SEQ ID
NO:26)] were used to amplify a 500-bp ***p53*** CDNA sequence by PCR
from the complete ***p53*** CDNA. This sequence was subcloned into
the pGEM-3zf(-) transcription vector (Promega Corp). The RNA probe was
used in northern blot. . . DETD

used in northern blot. . . . ID NO:27)] and another from the catalytic domain of the ribozyme sequence [5'-TCGTCCAAAAGGACTCATCAG-3' (SEQ ID NO:28)]. The level of endogenous ***p53*** expression was also assayed by RT

level of endogenous ***p53*** expression was also assayed by RT-PCR using a primer corresponding to ***exon*** 1 [5'-GGGAGAAAACGTTAGGGTGTG-3' (SEQ ID NO:29)] and ***exon*** 4 [5'-TGCAAGTCACAGACTTGGCTG-3' (SEQ ID NO:9)] of ***p53***. For northern blot analysis, the membrane was hybridized with an ***antisense*** ***p53*** RNA probe as described above. Hybridization and washing were performed according to the protocols supplied by Promega Corporation.

ANSWER 4 OF 32 USPATFULL L5

DETD

As demonstrated by the data shown herein and reported in Arora, 1998, the anti- ***p53*** ON suppress the expression of ***p53*** in post-hepatectomy rats. Morpholino and C-5-P ON's were found to be more effective than equivalent amounts of the unmodified phosphorothioate.

. liver. The morpholino oligomer is surprisingly effective in view of the fact that it is targeted at a coding region (***exon*** 10), well downstream of the AUG start region, the latter of which is the conventional target for RNAse-inactive ("steric blocker") ***antisense*** oligonucleotides. DETD